

GenCore version 5.1.6
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OM protein - protein search, using sw model

Run on: December 29, 2003, 14:19:12 ; Search time 643 Seconds
(without alignments)
181.375 Million cell updates/sec

Title: US-09-691-344A-4

Perfect score: 3052

Sequence: 1 MFGAGQRLRPVAPRSSAE.....YDNEKWTQKLDLITSDMAG 586

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 724715 seqs, 199017464 residues

Total number of hits satisfying chosen parameters: 724715

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : Published Applications AA:*

- 1: /cgn2_6/ptodata/1/pubpaa/PCTUS_PUBCOMB.pep:*
- 2: /cgn2_6/ptodata/1/pubpaa/PCTUS_PUBCOMB.pep:*
- 3: /cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*
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- 5: /cgn2_6/ptodata/1/pubpaa/US07_NEW_PUB.pep:*
- 6: /cgn2_6/ptodata/1/pubpaa/PCTUS_PUBCOMB.pep:*
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- 10: /cgn2_6/ptodata/1/pubpaa/US09B_PUBCOMB.pep:*
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- 13: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*
- 14: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*
- 15: /cgn2_6/ptodata/1/pubpaa/US10C_PUBCOMB.pep:*
- 16: /cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep:*
- 17: /cgn2_6/ptodata/1/pubpaa/US60_NEW_PUB.pep:*
- 18: /cgn2_6/ptodata/1/pubpaa/US60_PUBCOMB.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	3052	100.0	586	12	US-10-295-027-1300 Sequence 1300, Ap
2	2818	92.3	539	12	US-10-295-027-1299 Sequence 1299, Ap
3	2812	92.1	715	11	US-09-759-130B-73 Sequence 73, Appl
4	2812	92.1	715	14	US-10-003-132-2 Sequence 2, Appl
5	2652	86.9	681	11	US-09-759-130B-75 Sequence 75, Appl
6	2346.5	76.9	769	12	US-10-295-027-1301 Sequence 1301, Ap
7	2229	73.0	421	11	US-09-759-130B-76 Sequence 76, Appl
8	1309.5	42.9	503	14	US-10-003-132-4 Sequence 4, Appl
9	1301.5	42.6	503	9	US-09-823-038A-51 Sequence 51, Appl
10	1075	35.2	458	14	US-10-003-132-6 Sequence 6, Appl
11	942	30.9	769	16	US-10-191-436-8 Sequence 8, Appl
12	937	30.7	769	16	US-10-191-436-5 Sequence 5, Appl
13	929	30.4	775	16	US-10-191-436-2 Sequence 2, Appl
14	924.5	30.3	729	15	US-10-060-830-3 Sequence 3, Appl
15	906	29.7	554	12	US-10-094-749-2500 Sequence 2500, Ap

16	903.5	29.6	583	15	US-10-106-698-6224 Sequence 6224, Ap
17	756	24.8	653	15	US-10-060-830-1114 Sequence 1114, Ap
18	349	11.4	555	15	US-10-104-610-8 Sequence 8, Appl
19	349	11.4	926	12	US-10-247-671-171 Sequence 171, Appl
20	349	11.4	926	15	US-10-262-538-4 Sequence 4, Appl
21	349	11.4	931	14	US-10-104-440-4 Sequence 4, Appl
22	349	11.4	931	15	US-10-104-610-4 Sequence 4, Appl
23	343.5	11.3	926	15	US-10-262-538-8 Sequence 8, Appl
24	335	11.0	644	15	US-10-104-610-6 Sequence 6, Appl
25	335	11.0	923	14	US-10-104-440-2 Sequence 2, Appl
26	335	11.0	923	15	US-10-104-610-2 Sequence 2, Appl
27	335	11.0	923	15	US-10-262-538-2 Sequence 2, Appl
28	329	10.8	923	15	US-10-262-538-6 Sequence 6, Appl
29	317	10.4	2224	15	US-10-115-563-14 Sequence 14, Appl
30	317	10.4	2224	15	US-10-172-712-31 Sequence 31, Appl
31	307	10.1	235	11	US-09-759-130B-78 Sequence 78, Appl
32	306.5	10.0	1431	14	US-10-095-718-4 Sequence 4, Appl
33	299	9.8	343	15	US-10-190-593-2 Sequence 2, Appl
34	299	9.8	387	15	US-10-190-593-4 Sequence 4, Appl
35	285.5	9.4	1438	14	US-10-006-091-1 Sequence 1, Appl
36	285.5	9.4	1438	14	US-10-047-257-1 Sequence 1, Appl
37	285.5	9.4	1438	15	US-10-225-900-1 Sequence 2, Appl
38	285.5	9.4	1471	14	US-10-095-718-2 Sequence 2, Appl
39	285.5	9.4	2332	10	US-09-957-641-2 Sequence 2, Appl
40	285.5	9.4	2332	12	US-10-131-510A-2 Sequence 2, Appl
41	285.5	9.4	2332	15	US-10-187-319-2 Sequence 2, Appl
42	285.5	9.4	2351	12	US-10-133-907-4 Sequence 4, Appl
43	285.5	9.4	2351	15	US-10-132-829-4 Sequence 4, Appl
44	285.5	9.4	2351	15	US-10-172-712-27 Sequence 27, Appl
45	279.5	9.2	160	12	US-10-298-796-9 Sequence 9, Appl

ALIGNMENTS

RESULT 1

US-10-295-027-1300
; Sequence 1300, Application US/10295027
; Publication No. US20030232350A1
; GENERAL INFORMATION:
; APPLICANT: Afar, Daniel
; APPLICANT: Aziz, Natasha
; APPLICANT: Ginsberg, Wendy M.
; APPLICANT: Gish, Kurt C.
; APPLICANT: Glynn, Richard
; APPLICANT: Hevez, Peter A.
; APPLICANT: Mack, David H.
; APPLICANT: Murray, Richard
; APPLICANT: Watson, Susan R.
; APPLICANT: Eos Biotechnology, Inc.
; TITLE OF INVENTION: Methods of Diagnosis of Cancer, Compositions and Methods of Screening for Modulators of Cancer
; FILE OF INVENTION: 018501-012500US
; CURRENT APPLICATION NUMBER: US/10/295,027
; CURRENT FILING DATE: 2002-11-13
; PRIOR APPLICATION NUMBER: US 09/663,733
; PRIOR FILING DATE: 2000-09-15
; PRIOR APPLICATION NUMBER: US 60/350,666
; PRIOR FILING DATE: 2001-11-13
; PRIOR APPLICATION NUMBER: US 60/335,394
; PRIOR FILING DATE: 2001-11-15
; PRIOR APPLICATION NUMBER: US 60/332,464
; PRIOR FILING DATE: 2001-11-21
; PRIOR APPLICATION NUMBER: US 60/334,393
; PRIOR FILING DATE: 2001-11-29
; PRIOR APPLICATION NUMBER: US 60/340,376
; PRIOR FILING DATE: 2001-12-14
; PRIOR APPLICATION NUMBER: US 60/347,211
; PRIOR FILING DATE: 2002-01-08
; PRIOR APPLICATION NUMBER: US 60/347,349
; PRIOR FILING DATE: 2002-01-10
; PRIOR APPLICATION NUMBER: US 60/355,250
; PRIOR FILING DATE: 2002-02-08

; PRIOR APPLICATION NUMBER: US 60/356,714
; PRIOR FILING DATE: 2002-02-13
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 1386
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 1300
; LENGTH: 586
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-295-027-1300

Query Match 100.0%; Score 3052; DB 12; Length 586;
Best Local Similarity 100.0%; Pred. No. 1e-277;
Matches 586; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Db 61 AAGGGLALLAVSAPRLQAEELGDCGHLVTVQDSGTMTSKNYPGTYPNHTVCEKTI 120

Qy 121 VPKGKRLILRLGLDIDIESQTCASDYLLFTSSDQGYPCGSMTPVKELLNTSEVTVRFE 180
Db 121 VPKGKRLILRLGLDIDIESQTCASDYLLFTSSDQGYPCGSMTPVKELLNTSEVTVRFE 180

Qy 181 SGSHISGRGFLITVYASSDHPDLITCLERASHYLKTEYKPCPACGRDVAGDISGMVDGY 240
Db 181 SGSHISGRGFLITVYASSDHPDLITCLERASHYLKTEYKPCPACGRDVAGDISGMVDGY 240

Qy 241 RDTSLCKAAIHAGIIADELGGQISVLQKGISRYEGILANGVLSRDSLSDKRFLFTSN 300
Db 241 RDTSLCKAAIHAGIIADELGGQISVLQKGISRYEGILANGVLSRDSLSDKRFLFTSN 300

Qy 301 GCSRLSFPDQGIIRASSWSQVNESGDQVHWSFGQARLQDQGPSWASGDSNNHKPREW 360
Db 301 GCSRLSFPDQGIIRASSWSQVNESGDQVHWSFGQARLQDQGPSWASGDSNNHKPREW 360

Qy 361 LEIDLGEKKKITGRTTGSTOSNFFVYKSFVMNFKNNNSKWKTYKGIIVNEEKVFGNS 420
Db 361 LEIDLGEKKKITGRTTGSTOSNFFVYKSFVMNFKNNNSKWKTYKGIIVNEEKVFGNS 420

Qy 421 NFRDPVQNNFPIPIVARVVRVPTWHQRIALKVELIGCQITQGNDSLWKRKTSQTSVS 480
Db 421 NFRDPVQNNFPIPIVARVVRVPTWHQRIALKVELIGCQITQGNDSLWKRKTSQTSVS 480

Qy 481 TKKEDETITRPIPSEETSTGINITVAIPLVLLVLPAGMGIFAAPFRKKKKGSPYCSA 540
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Db 541 EAQKTDCKWKQIKYPFARHQSAEFTISYDNEKEMTKQLDLITSDMAG 586

RESULT 2

US-10-295-027-1299
; Sequence 1299, Application US/102995027
; Publication No. US2003023250A1
; GENERAL INFORMATION:
; APPLICANT: Afar, Daniel
; APPLICANT: Aziz, Natasha
; APPLICANT: Ginsberg, Wendy M.
; APPLICANT: Gish, Kurt C.
; APPLICANT: Glynn, Richard
; APPLICANT: Hevezi, Peter A.
; APPLICANT: Mack, David H.
; APPLICANT: Murray, Richard
; APPLICANT: Watson, Susan R.
; APPLICANT: Eos Biotechnology, Inc.
; TITLE OF INVENTION: Methods of Diagnosis of Cancer, Compositions and
; TITLE OF INVENTION: Methods of Screening for Modulators of Cancer

; FILE REFERENCE: 018501-012500US
; CURRENT APPLICATION NUMBER: US/10/295,027
; PRIOR FILING DATE: 2002-11-13
; PRIOR APPLICATION NUMBER: US 09/663,733
; PRIOR FILING DATE: 2000-09-15
; PRIOR APPLICATION NUMBER: US 60/350,666
; PRIOR FILING DATE: 2001-11-13
; PRIOR APPLICATION NUMBER: US 60/335,394
; PRIOR FILING DATE: 2001-11-15
; PRIOR APPLICATION NUMBER: US 60/332,464
; PRIOR FILING DATE: 2001-11-21
; PRIOR APPLICATION NUMBER: US 60/334,393
; PRIOR FILING DATE: 2001-11-29
; PRIOR APPLICATION NUMBER: US 60/340,376
; PRIOR FILING DATE: 2001-12-14
; PRIOR APPLICATION NUMBER: US 60/347,211
; PRIOR FILING DATE: 2002-01-08
; PRIOR APPLICATION NUMBER: US 60/347,349
; PRIOR FILING DATE: 2002-01-10
; PRIOR APPLICATION NUMBER: US 60/355,250
; PRIOR FILING DATE: 2002-02-08
; PRIOR APPLICATION NUMBER: US 60/356,714
; PRIOR FILING DATE: 2002-02-13
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 1386
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 1299
; LENGTH: 539
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-295-027-1299

Query Match 92.3%; Score 2818; DB 12; Length 539;
Best Local Similarity 100.0%; Pred. No. 8.9e-256;
Matches 539; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 48 MVFARGGGALARAAGRLALLAVSAPRLQAEELGDCGHLVTVQDSGTMTSKNYPG 107
Db 1 MVFARGGGALARAAGRLALLAVSAPRLQAEELGDCGHLVTVQDSGTMTSKNYPG 60

Qy 108 TYPNHTVCEKTIIVPKGRILILRLGLDIDIESQTCASDYLLFTSSDQGYPCGSMTPVKE 167
Db 61 TYPNHTVCEKTIIVPKGRILILRLGLDIDIESQTCASDYLLFTSSDQGYPCGSMTPVKE 120

Qy 168 LLNTSNTVTRFSGSHISGRGFLITVYASSDHPDLITCLERASHYLKTEYKPCPACGRD 227
Db 121 LLNTSNTVTRFSGSHISGRGFLITVYASSDHPDLITCLERASHYLKTEYKPCPACGRD 180

Qy 228 VAGDISGMVDGYRDTSLCKAAIHAGIIADELGGQISVLQKGISRYEGILANGVLSRD 287
Db 181 VAGDISGMVDGYRDTSLCKAAIHAGIIADELGGQISVLQKGISRYEGILANGVLSRD 240

Qy 288 GSLSDKRFLFTSNGCSRSLSEFPDQGIIRASSWSQVNESGDQVHWSFGQARLQDQGPSWA 347
Db 241 GSLSDKRFLFTSNGCSRSLSEFPDQGIIRASSWSQVNESGDQVHWSFGQARLQDQGPSWA 300

Qy 348 SGDSSNNHKPREWLEIDLGEKKKITGRTTGSTOSNFFVYKSFVMNFKNNNSKWKTYKG 407
Db 301 SGDSSNNHKPREWLEIDLGEKKKITGRTTGSTOSNFFVYKSFVMNFKNNNSKWKTYKG 360

Qy 408 IVNNEEKVFGNSNFRDPVQNNFPIPIVARVVRVPTWHQRIALKVELIGCQITQGNDS 467
Db 361 IVNNEEKVFGNSNFRDPVQNNFPIPIVARVVRVPTWHQRIALKVELIGCQITQGNDS 420

Qy 468 LWKRKTSQTSVSSTKKEDETIITRPIPSEETSTGINITVAIPLVLLVLPAGMGIFAAP 527
Db 421 LWKRKTSQTSVSSTKKEDETIITRPIPSEETSTGINITVAIPLVLLVLPAGMGIFAAP 480

Qy 528 RKKKKKGSPYCSAQAQKTDCKWKQIKYPFARHQSAEFTISYDNEKEMTKQLDLITSDMAG 586
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RESULT 3
US-09-759-130B-73
; Sequence 73, Application US/09/759130B
; Publication No. US20030022279A1
; GENERAL INFORMATION:
; APPLICANT: Millennium Pharmaceuticals, Inc.
; APPLICANT: McCarthy, Sean A
; APPLICANT: Fraser, Christopher C
; APPLICANT: Sharp, John D
; APPLICANT: Barnes, Thomas S
; APPLICANT: Kirst, Susan J
; APPLICANT: Mackay, Charles R
; APPLICANT: Myers, Paul S
; APPLICANT: Leiby, Kevin R
; APPLICANT: Wrighton, Nicolas
; APPLICANT: Goodearl, Andrew
; APPLICANT: Holtzman, Douglas A
; TITLE OF INVENTION: NOVEL GENES ENCODING PROTEINS HAVING
; TITLE OF INVENTION: PROGNOSTIC, DIAGNOSTIC, PREVENTIVE, THERAPEUTIC, AND OTHER
; TITLE OF INVENTION: USES.
; FILE REFERENCE: MPI00-5350MNI
; CURRENT APPLICATION NUMBER: US/09/759,130B
; CURRENT FILING DATE: 2002-09-16
; PRIOR APPLICATION NUMBER: US 09/479,249
; PRIOR FILING DATE: 2000-01-07
; PRIOR APPLICATION NUMBER: US 09/559,497
; PRIOR FILING DATE: 2000-04-27
; PRIOR APPLICATION NUMBER: US 09/578,063
; PRIOR FILING DATE: 2000-05-24
; PRIOR APPLICATION NUMBER: US 09/333,159
; PRIOR FILING DATE: 1999-06-14
; PRIOR APPLICATION NUMBER: US 09/596,194
; PRIOR FILING DATE: 2000-07-14
; PRIOR APPLICATION NUMBER: US 09/342,364
; PRIOR FILING DATE: 1999-06-29
; PRIOR APPLICATION NUMBER: US 09/608,452
; PRIOR FILING DATE: 2000-06-30
; PRIOR APPLICATION NUMBER: US 09/393,996
; PRIOR FILING DATE: 1999-09-10
; PRIOR APPLICATION NUMBER: US 09/602,871
; PRIOR FILING DATE: 2000-06-23
; PRIOR APPLICATION NUMBER: US 09/420,707
; PRIOR FILING DATE: 1999-10-19
; NUMBER OF SEQ ID NOS: 460
; SOFTWARE: Fast-Seq for Windows Version 4.0
; SEQ ID NO 73
; LENGTH: 715
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-759-130B-73

Query Match          92.1%; Score 2812; DB 11; Length 715;
Best Local Similarity 100.0%; Pred. No. 5.2e-255;
Matches 538; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Qy      108 T Y P N H T V C E K T I T V P K G R L I L R L G D L D I E S Q T C A S D Y L L F T S S D Q Y G P Y C G S M T V P K E 167
Db      61 T Y P N H T V C E K T I T V P K G R L I L R L G D L D I E S Q T C A S D Y L L F T S S D Q Y G P Y C G S M T V P K E 120

Qy      168 L L L N T S E V T V R F S G S H I S R G F L L T Y A S S D H P D L I T C L E R A S H Y L K T E Y S K F C P A G C R D 227
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Db      181 V A G D I S G N M V D G Y R D T S L L C K A A I H A G I I A D E L G G Q I S V L Q R K G I S R Y E G I L A N G V L S R D 240

Qy      288 G S L S D K R F L T S N G C S R S L S F E P D G Q I R A S S S W Q S V N E S G D Q V H W S P G Q A R L O D Q G P S W A 347
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Qy      348 S G D S S N N H K P R E W L E I D L G E K K I T G I R T T G S T Q S N F N F Y V K S F V M N F K N N S K W K T Y K G 407
Db      301 S G D S S N N H K P R E W L E I D L G E K K I T G I R T T G S T Q S N F N F Y V K S F V M N F K N N S K W K T Y K G 360

Qy      408 I V N N E K V F Q G N S N F R D P V Q N N F I P P I V A R Y V R V P Q T W H O R I A L K V E L I G C O I T G N D S 467
Db      361 I V N N E K V F Q G N S N F R D P V Q N N F I P P I V A R Y V R V P Q T W H O R I A L K V E L I G C O I T G N D S 420

Qy      467 I V N N E K V F Q G N S N F R D P V Q N N F I P P I V A R Y V R V P Q T W H O R I A L K V E L I G C O I T G N D S 467
Db      420 I V N N E K V F Q G N S N F R D P V Q N N F I P P I V A R Y V R V P Q T W H O R I A L K V E L I G C O I T G N D S 420
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Db      241 G S L S D K R F L T S N G C S R S L S F E P D G Q I R A S S S W Q S V N E S G D Q V H W S P G Q A R L O D Q G P S W A 300
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Db      301 S G D S S N N H K P R E W L E I D L G E K K I T G I R T T G S T Q S N F N F Y V K S F V M N F K N N S K W K T Y K G 360
Qy      408 I V N N E K V F Q G N S N F R D P V Q N N F I P P I V A R Y V R V P Q T W H O R I A L K V E L I G C O I T G N D S 467
Db      361 I V N N E K V F Q G N S N F R D P V Q N N F I P P I V A R Y V R V P Q T W H O R I A L K V E L I G C O I T G N D S 420
Qy      468 L V M R K T S Q S T S V T S K K E D E T I T R P I P S B E T S T G I N I T T V A I P L V L L V L V F A G M G I P A A F 527
Db      421 L V M R K T S Q S T S V T S K K E D E T I T R P I P S B E T S T G I N I T T V A I P L V L L V L V F A G M G I P A A F 480
Qy      528 R K K K K G S P Y G S A E A Q K T C W K Q I K Y P F A R H Q S A E F T I S Y D N E K E M T Q K L D L I T S D M A 585
Db      481 R K K K K G S P Y G S A E A Q K T C W K Q I K Y P F A R H Q S A E F T I S Y D N E K E M T Q K L D L I T S D M A 538

RESULT 4
US-10-003-132-2
; Sequence 2, Application US/10003132
; Publication No. US20020192750A1
; GENERAL INFORMATION:
; APPLICANT: Fox, Brian A.
; APPLICANT: Gao, Zeren
; APPLICANT: Shoemaker, Kimberly E.
; TITLE OF INVENTION: NEUROPTILIN HOMOLOG 2CUBS
; FILE REFERENCE: 00-62
; CURRENT APPLICATION NUMBER: US/10/003,132
; CURRENT FILING DATE: 2001-11-15
; PRIOR APPLICATION NUMBER: US 60/249,004
; PRIOR FILING DATE: 2000-11-15
; NUMBER OF SEQ ID NOS: 19
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 2
; LENGTH: 715
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-003-132-2

Query Match          92.1%; Score 2812; DB 14; Length 715;
Best Local Similarity 100.0%; Pred. No. 5.2e-255;
Matches 538; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy      48  M V P G A R G G A L A A R G L L A L L A V S A P L R L Q A E L G D G C G H L V T Y Q D S G T W T S K N Y P G 107
Db      1  M V P G A R G G A L A A R G L L A L L A V S A P L R L Q A E L G D G C G H L V T Y Q D S G T W T S K N Y P G 60

Qy      108 T Y P N H T V C E K T I T V P K G R L I L R L G D L D I E S Q T C A S D Y L L F T S S D Q Y G P Y C G S M T V P K E 167
Db      61 T Y P N H T V C E K T I T V P K G R L I L R L G D L D I E S Q T C A S D Y L L F T S S D Q Y G P Y C G S M T V P K E 120

Qy      168 L L L N T S E V T V R F S G S H I S R G F L L T Y A S S D H P D L I T C L E R A S H Y L K T E Y S K F C P A G C R D 227
Db      121 L L L N T S E V T V R F S G S H I S R G F L L T Y A S S D H P D L I T C L E R A S H Y L K T E Y S K F C P A G C R D 180

Qy      228 V A G D I S G N M V D G Y R D T S L L C K A A I H A G I I A D E L G G Q I S V L Q R K G I S R Y E G I L A N G V L S R D 287
Db      181 V A G D I S G N M V D G Y R D T S L L C K A A I H A G I I A D E L G G Q I S V L Q R K G I S R Y E G I L A N G V L S R D 240

Qy      288 G S L S D K R F L T S N G C S R S L S F E P D G Q I R A S S S W Q S V N E S G D Q V H W S P G Q A R L O D Q G P S W A 347
Db      241 G S L S D K R F L T S N G C S R S L S F E P D G Q I R A S S S W Q S V N E S G D Q V H W S P G Q A R L O D Q G P S W A 300

Qy      348 S G D S S N N H K P R E W L E I D L G E K K I T G I R T T G S T Q S N F N F Y V K S F V M N F K N N S K W K T Y K G 407
Db      301 S G D S S N N H K P R E W L E I D L G E K K I T G I R T T G S T Q S N F N F Y V K S F V M N F K N N S K W K T Y K G 360

Qy      408 I V N N E K V F Q G N S N F R D P V Q N N F I P P I V A R Y V R V P Q T W H O R I A L K V E L I G C O I T G N D S 467
Db      361 I V N N E K V F Q G N S N F R D P V Q N N F I P P I V A R Y V R V P Q T W H O R I A L K V E L I G C O I T G N D S 420
```

Qy 468 LWRKTSQSTSVSTKKEDETTIRIPSEETSTGINITTVAIPVLVLLVLFAGWIPAAAF 527
Db 421 LWRKTSQSTSVSTKKEDETTIRIPSEETSTGINITTVAIPVLVLLVLFAGWIPAAAF 480
Qy 528 RKKKKSPYGSABQAQKTCWQIKYFPARHQSNEFTISYDNEKEMTKQLDLITSDMA 585
Db 481 RKKKKSPYGSABQAQKTCWQIKYFPARHQSNEFTISYDNEKEMTKQLDLITSDMA 538
RESULT 5
US-09-759-130B-75
; Sequence 75, Application US/09759130B
; Publication No. US2003002279A1
; GENERAL INFORMATION:
; APPLICANT: Millennium Pharmaceuticals, Inc.
; APPLICANT: McCarthy, Sean A
; APPLICANT: Fraser, Christopher C
; APPLICANT: Sharp, John D
; APPLICANT: Barnes, Thomas S
; APPLICANT: Kirst, Susan J
; APPLICANT: Mackay, Charles R
; APPLICANT: Myers, Paul S
; APPLICANT: Leiby, Kevin R
; APPLICANT: Wrighton, Nicolas
; APPLICANT: Goodearl, Andrew
; APPLICANT: Holtzman, Douglas A
; TITLE OF INVENTION: NOVEL GENES ENCODING PROTEINS HAVING
; TITLE OF INVENTION: PROGNOSTIC, DIAGNOSTIC, PREVENTIVE, THERAPEUTIC, AND OTHER
; TITLE OF INVENTION: USES
; FILE REFERENCE: MP100-535OMNIM
; CURRENT APPLICATION NUMBER: US/09/759,130B
; CURRENT FILING DATE: 2002-09-16
; PRIOR APPLICATION NUMBER: US 09/479,249
; PRIOR FILING DATE: 2000-01-07
; PRIOR APPLICATION NUMBER: US 09/559,497
; PRIOR FILING DATE: 2000-04-27
; PRIOR APPLICATION NUMBER: US 09/578,063
; PRIOR FILING DATE: 2000-05-24
; PRIOR APPLICATION NUMBER: US 09/333,159
; PRIOR FILING DATE: 1999-06-14
; PRIOR APPLICATION NUMBER: US 09/596,194
; PRIOR FILING DATE: 2000-07-14
; PRIOR APPLICATION NUMBER: US 09/342,364
; PRIOR FILING DATE: 1999-06-29
; PRIOR APPLICATION NUMBER: US 09/608,452
; PRIOR FILING DATE: 2000-06-30
; PRIOR APPLICATION NUMBER: US 09/393,996
; PRIOR FILING DATE: 1999-09-10
; PRIOR APPLICATION NUMBER: US 09/602,871
; PRIOR FILING DATE: 2000-06-23
; PRIOR APPLICATION NUMBER: US 09/420,707
; PRIOR FILING DATE: 1999-10-19
; NUMBER OF SEQ ID NOS: 460
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 75
; LENGTH: 681
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-759-130B-75
Query Match 86.9%; Score 2652; DB 11; Length 681;
Best Local Similarity 100.0%; Pred. No. 5.3e-240;
Matches 504; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
Qy 82 BELGGCGHLVYQDSGMTSKNYPGYNHTVCEKTIYVPGKRLILRLGLDLIESQTC 141
Db 1 BELGGCGHLVYQDSGMTSKNYPGYNHTVCEKTIYVPGKRLILRLGLDLIESQTC 60
Qy 142 ASDVLLFTSSSDQGPYCGSMVTVPKELLNTSEVTVPFSGSHISGRGFLITYASSDHPD 201
Db 61 ASDVLLFTSSSDQGPYCGSMVTVPKELLNTSEVTVPFSGSHISGRGFLITYASSDHPD 120
Qy 202 LITCLERASHYLKTEYSKFCPAGCRDVAGDISGNMVDGYRDTSLCKAAIHAGIIADELG 261

Db 121 LITCLERASHYLKTEYSKFCPAGCRDVAGDISGNMVDGYRDTSLCKAAIHAGIIADELG 180
Qy 262 GOISVLQRKIGISRYEGILANGVLSRDGSLDKRFLFTSNGCSRSLSFEPDGOIRASSSQ 321
Db 181 GOISVLQRKIGISRYEGILANGVLSRDGSLDKRFLFTSNGCSRSLSFEPDGOIRASSSQ 240
Qy 322 SVNESGDQVHWSFGQARLQDQGPSWASGDSNNHKPREWLEIDLGEKKKITGIRTTGSTQ 381
Db 241 SVNESGDQVHWSFGQARLQDQGPSWASGDSNNHKPREWLEIDLGEKKKITGIRTTGSTQ 300
Qy 382 SNPNFYVKSFMNFKNNNSKWKTYKGI VNNNEKVFQGNFNFRDPVQNNFIPPIVARYVRV 441
Db 301 SNPNFYVKSFMNFKNNNSKWKTYKGI VNNNEKVFQGNFNFRDPVQNNFIPPIVARYVRV 360
Qy 442 VPQTHORIALKVELIGCOITQGNDSLVRKTSQSTSVSTKKEDETTIRIPSEETSTGI 501
Db 361 VPQTHORIALKVELIGCOITQGNDSLVRKTSQSTSVSTKKEDETTIRIPSEETSTGI 420
Qy 502 NITTVAILPLVLVLFAGWIFAAFRKKKKGSPYGSABQAQKTCWQIKYFPARHQA 561
Db 421 NITTVAILPLVLVLFAGWIFAAFRKKKKGSPYGSABQAQKTCWQIKYFPARHQA 480
Qy 562 EFTISYDNEKEMTKQLDLITSDMA 585
Db 481 EFTISYDNEKEMTKQLDLITSDMA 504
RESULT 6
US-10-295-027-1301
; Sequence 1301, Application US/10295027
; Publication No. US20030232350A1
; GENERAL INFORMATION:
; APPLICANT: Afar, Daniel
; APPLICANT: Aziz, Natasha
; APPLICANT: Ginsberg, Wendy M.
; APPLICANT: Gish, Kurt C.
; APPLICANT: Glynn, Richard
; APPLICANT: Hevezi, Peter A.
; APPLICANT: Mack, David H.
; APPLICANT: Murray, Richard
; APPLICANT: Watson, Susan R.
; APPLICANT: Eos Biotechnology, Inc.
; TITLE OF INVENTION: Methods of Diagnosis of Cancer, Compositions and
; FILE REFERENCE: 018501-012500US
; CURRENT APPLICATION NUMBER: US/10/295,027
; CURRENT FILING DATE: 2002-11-13
; PRIOR APPLICATION NUMBER: US 09/663,733
; PRIOR FILING DATE: 2000-09-15
; PRIOR APPLICATION NUMBER: US 60/350,666
; PRIOR FILING DATE: 2001-11-13
; PRIOR APPLICATION NUMBER: US 60/335,394
; PRIOR FILING DATE: 2001-11-15
; PRIOR APPLICATION NUMBER: US 60/332,464
; PRIOR FILING DATE: 2001-11-21
; PRIOR APPLICATION NUMBER: US 60/334,393
; PRIOR FILING DATE: 2001-11-29
; PRIOR APPLICATION NUMBER: US 60/340,376
; PRIOR FILING DATE: 2001-12-14
; PRIOR APPLICATION NUMBER: US 60/347,211
; PRIOR FILING DATE: 2002-01-08
; PRIOR APPLICATION NUMBER: US 60/347,349
; PRIOR FILING DATE: 2002-01-10
; PRIOR APPLICATION NUMBER: US 60/355,250
; PRIOR FILING DATE: 2002-02-08
; PRIOR APPLICATION NUMBER: US 60/356,714
; PRIOR FILING DATE: 2002-02-13
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 1386
; SOFTWARE: Patent in Ver. 2.1
; SEQ ID NO 1301
; LENGTH: 769

```
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-295-027-1301

Query Match      76.9%; Score 2346.5; DB 12; Length 769;
Best Local Similarity 78.7%; Pred. No. 3.4e-211;
Matches 466; Conservative 13; Mismatches 20; Indels 93; Gaps 6;

Qy 72 AVSAPRLQABEL---GDGCGHLVTVQDSGTMTSKNYPGTYPNHTVCEKITVPKGRLLI 128
Db : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Qy 16 AVAAWILHQHDDIINKGDCGHLVTVQDSGTMTSKNYPGTYPNHTVCEKITVPKGRLLI 75
Db : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Qy 129 LRLGDLIESQTCADYLLFTSSSDQY----- 155
Db : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Qy 76 LRLGDLIESQTCADYLLFTSSSDQYCMQKEETEVLCLSVAGAQRVDPVQLLPSPLE 135
Db : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Qy 156 -----GPYCGSMVTPKELLANTSEVTVRFESGSHISGRGFLLYASSDHPDLITCL 206
Db : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Qy 136 GWKGHADARGPYCGSMVTPKELLANTSEVTVRFESGSHISGRGFLLYASSDHPDLITCL 195
Db : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Qy 207 ERASHYLKTEYSKFCPCAGCRDVAGDISGNMVDGYRDTSLCKAAIHAGIIADELGGQISV 266
Db : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Qy 196 ERASHYLKTEYSKFCPCAGCRDVAGDISGNMVDGYRDTSLCKAAIHAGIIADELGGQISV 255
Db : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Qy 267 LQKIGISRYEGILANGVLSRDGSLSDKRFLLTSNGCSRSLSEFPDQIRASSSSWQSVNES 326
Db : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Qy 256 LQKIGISRYEGILANGVLSRDGSLSDKRFLLTSNGCSRSLSEFPDQIRASSSSWQSVNES 315
Db : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Qy 327 GDQVHWSFGQARLODQGPSWASGSSNNHKPREWLEIDLGEKKITGIRTTGTSQSNFN 386
Db : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Qy 316 GDQVHWSFGQARLODQGPSWASGSSNNHKPREWLEIDLGEKKITGIRTTGTSQSNFN 375
Db : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Qy 387 YVKSFVMNFKNNNSKWKTYKGI VVNEEKVFCGNSNFRDPVQNNFIPPIVARYVRVPTW 446
Db : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Qy 376 YVKSFVMNFKNNNSKWKTYKGI VVNEEKVFCGNSNFRDPVQNNFIPPIVARYVRVPTW 435
Db : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Qy 447 HQRIALKVELIGCQITQGNDSLVRKTSQSSTSVTKKEDETITRPIPEETSTGINITTV 506
Db : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Qy 436 HQRIALKVELIGCQITQGNDSLVRKTSQSSTSVTKKEDETITRPIPEETSTGTD----- 489
Db : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Qy 507 AIPVLV-----LVLVFA-----GMGFAAFKKKK 533
Db : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Qy 490 AMPQIVGDHTQMTISQRENLPDGGKIPFKGTASVMRVVFAVVNDLGMFLAHTPEED 549
Db : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Qy 534 GSPYGSAAEQTKDCWKQIKYFARHQSAEFTISVDNEKEMTKQLDLITSDMA 585
Db : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Qy 550 IDHY-----CWKQIKYFARHQSAEFTISVDNEKEMTKQLDLITSDMA 592
Db : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :

RESULT 7
US-09-759-130B-76
; Sequence 76, Application US/09759130B
; Publication No. US20030022279A1
; GENERAL INFORMATION:
; APPLICANT: Millennium Pharmaceuticals, Inc.
; APPLICANT: McCarthy, Sean A
; APPLICANT: Fraser, Christopher C
; APPLICANT: Sharp, John D
; APPLICANT: Barnes, Thomas S
; APPLICANT: Kirt, Susan J
; APPLICANT: Mackay, Charles R
; APPLICANT: Myers, Paul S
; APPLICANT: Leiby, Kevin R
; APPLICANT: Wrighton, Nicolas
; APPLICANT: Goodearl, Andrew
; APPLICANT: Holtzman, Douglas A
; TITLE OF INVENTION: NOVEL GENES ENCODING PROTEINS HAVING
; TITLE OF INVENTION: PROGNOSTIC, DIAGNOSTIC, PREVENTIVE, THERAPEUTIC, AND OTHER
; TITLE OF INVENTION: USES
; FILE REFERENCE: MP100-535OMNIM
; CURRENT APPLICATION NUMBER: US/09/759,130B
; CURRENT FILING DATE: 2002-09-16
; PRIOR APPLICATION NUMBER: US 09/479,249

; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-295-027-1301

Query Match      73.0%; Score 2229; DB 11; Length 421;
Best Local Similarity 100.0%; Pred. No. 1.4e-200;
Matches 421; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 82 EELGDCGCHLVTVQDSGTMTSKNYPGTYPNHTVCEKITVPKGRLLIRLGLDLIESQTC 141
Db : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Qy 142 ASDYLLFTSSSDQYGPYCGSMVTPKELLANTSEVTVRFESGSHISGRGFLLYASSDHPD 201
Db : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Qy 61 ASDYLLFTSSSDQYGPYCGSMVTPKELLANTSEVTVRFESGSHISGRGFLLYASSDHPD 120
Db : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Qy 202 LITCLERASHYLKTEYSKFCPCAGCRDVAGDISGNMVDGYRDTSLCKAAIHAGIIADEL 261
Db : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Qy 121 LITCLERASHYLKTEYSKFCPCAGCRDVAGDISGNMVDGYRDTSLCKAAIHAGIIADEL 180
Db : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Qy 262 GOISVLQKIGISRYEGILANGVLSRDGSLSDKRFLLTSNGCSRSLSEFPDQIRASSSWQ 321
Db : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Qy 181 GOISVLQKIGISRYEGILANGVLSRDGSLSDKRFLLTSNGCSRSLSEFPDQIRASSSWQ 240
Db : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Qy 322 SVNESGDQVHWSFGQARLODQGPSWASGSSNNHKPREWLEIDLGEKKITGIRTTGTSQ 381
Db : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Qy 241 SVNESGDQVHWSFGQARLODQGPSWASGSSNNHKPREWLEIDLGEKKITGIRTTGTSQ 300
Db : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Qy 382 SNFNFYVKSFVMNFKNNNSKWKTYKGI VVNEEKVFCGNSNFRDPVQNNFIPPIVARYVRV 441
Db : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Qy 301 SNFNFYVKSFVMNFKNNNSKWKTYKGI VVNEEKVFCGNSNFRDPVQNNFIPPIVARYVRV 360
Db : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Qy 442 VPQTHORIALKVELIGCQITQGNDSLVRKTSQSSTSVTKKEDETITRPIPEETSTGI 501
Db : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Qy 361 VPQTHORIALKVELIGCQITQGNDSLVRKTSQSSTSVTKKEDETITRPIPEETSTGI 420
Db : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Qy 502 N 502
Db : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Qy 421 N 421
Db : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :

RESULT 8
US-10-003-132-4
; Sequence 4, Application US/10003132
; Publication No. US20020192750A1
; GENERAL INFORMATION:
; APPLICANT: Fox, Brian A.
; APPLICANT: Gao, Zeren
; APPLICANT: Shoemaker, Kimberly E.
; TITLE OF INVENTION: NEUROFILIN HOMOLOG ZCUB5
```

```
; FILE REFERENCE: 00-62
; CURRENT APPLICATION NUMBER: US/10/003.132
; CURRENT FILING DATE: 2001-11-15
; PRIOR APPLICATION NUMBER: US 60/249,004
; PRIOR FILING DATE: 2000-11-15
; NUMBER OF SEQ ID NOS: 19
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 4
; LENGTH: 503
; TYPE: PRT
; ORGANISM: Mus musculus
; US-10-003-132-4

Query Match      42.9%; Score 1309.5; DB 14; Length 503;
Best Local Similarity 51.0%; Pred. No. 5.5e-114;
Matches 268; Conservative 26; Mismatches 32; Indels 199; Gaps 1;

Qy 61 AAGRGLLALLAVSAPLRQAELGDCGHLVTVQDSGTMTSKNYPGTYPNHTVCEKTIIT 120
Db 5 AGGPSVLALLFAVCAPLRQAELGDCGHLVTVQDSGTMTSKNYPGTYPNHTVCEKTIIT 64

Qy 121 VPKGKRLILRLGLDLDIESOTCASDYLLFTSSSDQYGPYCGSMTVPKELLINTSEVTVRFE 180
Db 65 VPKGKRLILRLGLDLDIESOTCASDYLLFTSSSDQYGPYCGSMTVPKELLINTSEVTVRFE 124

Qy 181 SGSHISGRGFLTYASSDHPDLITCLERASHYLKTEYKFCPCAGCRDVAGDISGNMVDGY 240
Db 125 SGSHISGRGFLTYASSDHPDLITCLERASHYLKTEYKFCPCAGCRDVAGDISGNMVDGY 184

Qy 241 RDTSLCKAAIHAGIITDELGGQISVLQKGISRYEGILANGVLSRDGSLDKRFLFTSN 300
Db 185 RDTSLCKAAIHAGIITDELGGHINLLQSKGISHYEGLLANGVLSRHGSLSEKRFLE--- 241

Qy 301 GCSRSLSFEPDQGIIRASSWSQSVNESGDQVHWSFGQARLQDQGSWASGDSNNHKKPREW 360
Db 242 ----- 241

Qy 361 LEIDLGEKKKITGIRTTGSTQSNFNFYKSPVMFNKNNNSKWKTYKGINNEEKVFGNS 420
Db 242 ----- 241

Qy 421 NFRDPVQNNFPPIVARYVRVPQTHQRIALKVELIGCQITQGNDSLWRKTSQSTSUS 480
Db 242 ----- 241

Qy 481 TKKEDETITRIPSEETSTGINITTVAIPVLVLVLFAGMGIFAAPRKKKKKSPYGSA 540
Db 242 -----TTPGMNITTVAIPSVIFIALLTGGMGIFAICRKKKGNPYVSA 285

Qy 541 EAQKTCWKQIKYPPFARHOSAEFTISYDNEKEMTKQLDLITSDMA 585
Db 286 DAQKTCWKQIKYPPFARHOSAEFTISYDNEKEMTKQLDLITSDMA 330

RESULT 9
US-09-823-038A-51
; Sequence 51, Application US/09823038A
; Patent No. US20020058335A1
; GENERAL INFORMATION:
; APPLICANT: Strachan, Lorna
; APPLICANT: Sleeman, Matthew
; APPLICANT: Abernethy, Nevin
; APPLICANT: Onrust, Rene
; APPLICANT: Kumble, Anand
; APPLICANT: Murison, Greg
; TITLE OF INVENTION: Compositions Isolated From Stromal Cells
; FILE REFERENCE: 11000.1037c3
; CURRENT APPLICATION NUMBER: US/09/823,038A
; CURRENT FILING DATE: 2001-07-09
; NUMBER OF SEQ ID NOS: 61
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 51

; FILE REFERENCE: 00-62
; CURRENT APPLICATION NUMBER: US/10/003.132
; CURRENT FILING DATE: 2001-11-15
; PRIOR APPLICATION NUMBER: US 60/249,004
; PRIOR FILING DATE: 2000-11-15
; NUMBER OF SEQ ID NOS: 19
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 4
; LENGTH: 503
; TYPE: PRT
; ORGANISM: Mus musculus
; US-10-003-132-6

Query Match      42.6%; Score 1301.5; DB 9; Length 503;
Best Local Similarity 50.9%; Pred. No. 3.1e-113;
Matches 267; Conservative 26; Mismatches 33; Indels 199; Gaps 1;

Qy 61 AAGRGLLALLAVSAPLRQAELGDCGHLVTVQDSGTMTSKNYPGTYPNHTVCEKTIIT 120
Db 5 AGGPSVLALLFAVCAPLRQAELGDCGHLVTVQDSGTMTSKNYPGTYPNHTVCEKTIIT 64

Qy 121 VPKGKRLILRLGLDLDIESOTCASDYLLFTSSSDQYGPYCGSMTVPKELLINTSEVTVRFE 180
Db 65 VPKGKRLILRLGLDLDIESOTCASDYLLFTSSSDQYGPYCGSMTVPKELLINTSEVTVRFE 124

Qy 181 SGSHISGRGFLTYASSDHPDLITCLERASHYLKTEYKFCPCAGCRDVAGDISGNMVDGY 240
Db 125 SGSHISGRGFLTYASSDHPDLITCLERASHYLKTEYKFCPCAGCRDVAGDISGNMVDGY 184

Qy 241 RDTSLCKAAIHAGIITDELGGQISVLQKGISRYEGILANGVLSRDGSLDKRFLFTSN 300
Db 185 RDTSLCKAAIHAGIITDELGGHINLLQSKGISHYEGLLANGVLSRHGSLSEKRFLE--- 241

Qy 301 GCSRSLSFEPDQGIIRASSWSQSVNESGDQVHWSFGQARLQDQGSWASGDSNNHKKPREW 360
Db 242 ----- 241

Qy 361 LEIDLGEKKKITGIRTTGSTQSNFNFYKSPVMFNKNNNSKWKTYKGINNEEKVFGNS 420
Db 242 ----- 241

Qy 421 NFRDPVQNNFPPIVARYVRVPQTHQRIALKVELIGCQITQGNDSLWRKTSQSTSUS 480
Db 242 ----- 241

Qy 481 TKKEDETITRIPSEETSTGINITTVAIPVLVLVLFAGMGIFAAPRKKKKKSPYGSA 540
Db 242 -----TTPGMNITTVAIPSVIFIALLTGGMGIFAICRKKKGNPYVSA 285

Qy 541 EAQKTCWKQIKYPPFARHOSAEFTISYDNEKEMTKQLDLITSDMA 585
Db 286 DAQKTCWKQIKYPPFARHOSAEFTISYDNEKEMTKQLDLITSDMA 330

RESULT 10
US-10-003-132-6
; Sequence 6, Application US/10003132
; Publication No. US20020192750A1
; GENERAL INFORMATION:
; APPLICANT: Fox, Brian A.
; APPLICANT: Gao, Zeren
; APPLICANT: Shoemaker, Kimberly E.
; TITLE OF INVENTION: NEUROFILIN HOMOLOG ZCUB5
; FILE REFERENCE: 00-62
; CURRENT APPLICATION NUMBER: US/10/003.132
; CURRENT FILING DATE: 2001-11-15
; PRIOR APPLICATION NUMBER: US 60/249,004
; PRIOR FILING DATE: 2000-11-15
; NUMBER OF SEQ ID NOS: 19
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 6
; LENGTH: 458
; TYPE: PRT
; ORGANISM: Mus musculus
; US-10-003-132-6

Query Match      35.2%; Score 1075; DB 14; Length 458;
Best Local Similarity 43.8%; Pred. No. 5.3e-92;
Matches 230; Conservative 23; Mismatches 28; Indels 244; Gaps 2;

Qy 61 AAGRGLLALLAVSAPLRQAELGDCGHLVTVQDSGTMTSKNYPGTYPNHTVCEKTIIT 120
```

Db 5 AGGSLVALLPAVCAPIRLQAEEIGDGGHIVTSQDSGTMTSKNYPGTYNVTCEKIIT 64
Qy 121 VPKGRILRLGLDIDIESQTCASDYLLFTSSDQYGPYCGSMTVPKELLNLTSEVTRFE 180
Db 65 VPKGRILRLGLDNLIESKTCASDYLLFSSATDQY----- 99
Qy 181 SGSHISGRGFLITYASSDHPDLITCLERASHYLKTEYSKFCPACGRDVAGDISGNMVDGY 240
Db 100 -----DLITCLERGSHYFEKYSKFCPACGRDIAGDISGNTKGY 139
Qy 241 RDTSLCKAAHAGIIADELGGQISVLRKGISRYEGILANGVLSRDGSLSDKRFLETSN 300
Db 140 RDTSLCKAAHAGIITDELGGINLQSGKISHYEGILANGVLSRHGSLSEKRFLEP--- 196
Qy 301 GCSRSLSFEPDGOIRASSWSQVNESGDQVHWSPGQARLQDQGSWASGDSNNHKKREW 360
Db 197 ----- 196
Qy 361 LEIDLGKKKITGIRTTGTSQSNFNFYKGFVMNFKNNKWKTYKGIIVNNEEKVFGQNS 420
Db 197 ----- 196
Qy 421 NFRDPQNNFIPPIVARYVRVQVTHQRIALKVELIGCOITQGNDSLVRKTSQTSVS 480
Db 197 ----- 196
Qy 481 TKKEDETIRIPSEETSTGINITVAIPLVLLVLFVAGMGIFAAPFRKKKXGSPYGSA 540
Db 197 -----TTPGNITVAIPSVIFIALLTGWIFAICRKRKKNPIYVA 240
Qy 541 EAQKTCWKQIKYPPFARHQAFTISYDNKEMTKQLDLITSDMA 585
Db 241 DAQKTCWKQIKYPPFARHQSTFTISYDNKEMTKQLDLITSDMA 285

RESULT 11

US-10-191-436-8
; Sequence 8, Application US/10191436
; Publication No. US20030129697A1
; GENERAL INFORMATION:
; APPLICANT: Tasuku Honjo
; APPLICANT: Kei Tashiro
; APPLICANT: Kazuhiro Kobuke
; TITLE OF INVENTION: A NOVEL POLYPEPTIDE ESDN, POLYNUCLEOTIDES ENCODING THE POLYPEPTIDE
; FILE REFERENCE: ONF3204US
; CURRENT FILING DATE: 2002-07-10
; CURRENT APPLICATION NUMBER: US/10/191,436
; NUMBER OF SEQ ID NOS: 24
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 8
; LENGTH: 769
; TYPE: PRT
; ORGANISM: Rattus rattus
US-10-191-436-8

Query Match 30.9%; Score 942; DB 16; Length 769;
Best Local Similarity 37.5%; Pred. No. 3.9e-79;
Matches 230; Conservative 96; Mismatches 204; Indels 84; Gaps 19;

Qy 44 PSGVMVPGARGGGALAAA-----GRG-----LLALLAVSAPRLQAELGDCG 89
Db 16 PGGRAAPAATGRAPLSAGWCPLPPGGRNSSRRPRLLLLLLLLPDA-----GAQKGDGCG 70
Qy 90 HLYTQDSGTMTSKNYPGTYNHTVCEKTIIVPKGRILRLGLDIE-SQTCASDYLL-L 147
Db 71 HTVLGPESGLTISINPHYTPNSTVCEWEIRVKTGERIRIKFGDFDIEDSDYCHLVYKI 130
Qy 148 FTS---SSDQGYPCG-SMTVPKELLNLTSEVTRFESGSHISGRGFLITYASSDHPDLI 203
Db 131 FNGIGVSRTEIGKYCGLGLQMNQSIKSGSEITVLFMSGHASGRGLFASYSVIDKODLI 190

Qy 204 TCLERASHYLKTEYSKFCPACGRDVAGDISGNMVDGYRDTSLCKAAHAGIIADELGGQ 263
Db 191 TCLDTVSNFLEPEFSKYCPAGCLLPFAEISGTIPHGVRDSSPLCMAGIHAGVVDVLGGQ 250
Qy 264 ISVLQRKGISRYEGILANGVLSRDGSLSDKRFLETSNGCSRSLSFE-----PDQIRASS 319
Db 251 ISVVISKGTPTPYBSSLANNVTSVMGYLSTLSLFTFKTSGCYTILGMEGVIADPQITASSV 310
Qy 320 WQSVNESGDQVHWSPGQARLQDQGSWASGDSNNHKKREWLEIDLCEKKKITGIRTTGS 379
Db 311 LEWTDHMGQENSKWPEKARLKFGPPWAA-FATDEH---QWLQIDLNKEKKITGIVTTGS 366
Qy 380 TQSNFNFYKGFVMNFKNNKWKTYKGIIVNNEEKVFGQNSFRDPVQNNFIPPIVARYV 439
Db 367 TLIENHYVYSAYRVLSDDGQKWTVYREPCGAADKIFQGNKYDKVDRNNFLPPIARFI 426
Qy 440 RVVPQVTHQRIALKVELIGCOIT-----QGNDSLVRKTSOSTSVSTYKED 485
Db 427 RVNPFVQWQQKIAMKVELLGCQFTLKGRLPKLTOPPPRNSNL--KNTTVHPLKGRAPKF 484
Qy 486 ETITRP-----IPSEETST-GINITT-----VAIPLVLLVLFVAGMGIFAAP- 527
Db 485 TQALQPRSRNDLPLPAQTATPDVKNVTTPSVTKDVAALVLPVLVWALTLLILV 544
Qy 528 -----RKKKKXGSPYSAEAQKTCWKQIK--YPPFARHQAFTISYDNE-----K 571
Db 545 CAWHNRNKKKABGT-YDLPHMDRAGWKGVKQLLPAKSVEHEETPRYSNSEVSHLSPR 603
Qy 572 EMTQKLDLITSDMA 585
Db 604 EVTTVLQADSAEA 617

RESULT 12

US-10-191-436-5
; Sequence 5, Application US/10191436
; Publication No. US20030129697A1
; GENERAL INFORMATION:
; APPLICANT: Tasuku Honjo
; APPLICANT: Kei Tashiro
; APPLICANT: Kazuhiro Kobuke
; TITLE OF INVENTION: A NOVEL POLYPEPTIDE ESDN, POLYNUCLEOTIDES ENCODING THE POLYPEPTIDE
; FILE REFERENCE: ONF3204US
; CURRENT FILING DATE: 2002-07-10
; CURRENT APPLICATION NUMBER: US/10/191,436
; NUMBER OF SEQ ID NOS: 24
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 5
; LENGTH: 769
; TYPE: PRT
; ORGANISM: Mus musculus
US-10-191-436-5

Query Match 30.7%; Score 937; DB 16; Length 769;
Best Local Similarity 37.2%; Pred. No. 1.2e-78;
Matches 229; Conservative 100; Mismatches 200; Indels 86; Gaps 17;

Qy 44 PSGVMVPGARGGGALAAA-----GRG-----LLALLAVSAPRLQAELGDCGHL 91
Db 16 PGGAAPAATGRAALPSAGCCPLPPGGRNSSRRPRLLLLLLL---LQDAGGQGGCGHT 72
Qy 92 VTYQDSGTMTSKNYPGTYNHTVCEKTIIVPKGRILRLGLDIE-SQTCASDYLL-LFT 149
Db 73 VLGPESGLTISINPHYTPNSTVCEWEIRVKTGERIRIKFGDFDIEDSDYCHLVYKI 132
Qy 150 S---SSDQGYPCG-SMTVPKELLNLTSEVTRFESGSHISGRGFLITYASSDHPDLITC 205
Db 133 GIGVSRTEIGKYCGLGLQMNQSIKSGSEITVLFMSGHASGRGLFASVIDKEDLIIC 192
Qy 206 LERASHYLKTEYSKFCPACGRDVAGDISGNMVDGYRDTSLCKAAHAGIIADELGGQIS 265
Db 193 LDTVSNFLEPEFSKYCPAGCLLPFAEISGTIPHGVRDSSPLCMAGIHAGVVDVLGGQIS 252


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181  SGSHISGRGFLITYASSDPDLITCLERASHYHLTEYSKFCPCAGCDVAGDISGNMVDGY 240
182  ||:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:
125  SGIHVSGRGFLIASSVIDKDLITCLDTASNFLEPEFSKYCPAGCLLPFAEISGTHIPGHY 184
186  ||:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:
241  RDTSLCLKAAIHAGIADBLGGQISVLORKGISRYEGILANGVLSDGSLSKDRFLFTSN 300
188  ||:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:
185  RDSFPLCWAGHVAGVSNLTGGQISVWISKGIPIYYESSLANNVTSVWGHLSLSTFFKTS 244
189  ||:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:
301  GCSSLSLFE---PDGQIRASSWQSVNESGDVHWSPCQARLODQPSWASGDSNNHK 356
191  ||:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:
245  GCYGTGLWGESVIADPQITASVLEWTHTCQENSWKPKAKLKPGPPWPAFAFDE--- 301
193  ||:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:
357  PREWLEIDLGBKKKITGIRTTGSTQSNFNFYVKSFWMNFKNNSKWKTYKGINNEEKVF 416
195  ||:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:
302  -YQWLQDLNLNKEKKITGIITGSTVMEHNYVSAYRILYSDGQKWTVYREPGVEQDKIF 360
197  ||:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:
417  QGNSNFRDPVONNTPPIVARYVRVPQTHQRIALKVELIGCQ-----ITQ----- 463
199  ||:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:
361  QGNKDYHQDVNNFLPPIIAFIRVNPQWQOKIAMKMLGCCQIPKGRPPKLIQPPPP 420
201  ||:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:
464  --GNDLSLWRKTSOSTSVSTKKEDEITRPI-----PSEBT--STGINIIT----- 505
203  ||:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:
421  RNSND-----LKNTPAPPRIAKGRAPKFTQPLQPRSSNEFPQTEQTASPDIRNTTVTPN 476
205  ||:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:
506  ---VAIPVLVLVLVAFAGMGIFAFAF-----RKKKKKS---PYGSAEAQKTDCKWQ 550
207  ||:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:
477  VTKDVALAANVLVPVLVMVLTLLILLCVAWHNRNKKKTEGYIDLPYW---DRAGWK 532
209  ||:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:
551  IKYPF-----ARHQSAEFTISYDNBKEMTQKLDLITSOWA 585
211  ||:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:
533  MKQPLPAKADVHEETPVRYSSSE---VNHLSPREVTVLQADSAEA 576
213  ||:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:|||||:|:

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RESULT 15

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US-10-094-749-2500
; Sequence 2500, Application US/10094749
; Publication No. US20030219741A1
; GENERAL INFORMATION:
; APPLICANT: ISOGAI, TAKAO
; APPLICANT: SUGIYAMA, TOMOYASU
; APPLICANT: OTSUKI, TETSUJI
; APPLICANT: WAKAMATSU, AI
; APPLICANT: SATO, HIROYUKI
; APPLICANT: ISHII, SHIZUKU
; APPLICANT: YAMAMOTO, JUN-ICHI
; APPLICANT: ISONO, YUUKO
; APPLICANT: HIO, YURI
; APPLICANT: OTSUKA, KAORU
; APPLICANT: NAGAI, KEIICHI
; APPLICANT: IRIE, RYOTARO
; APPLICANT: TAMECHIKA, ICHIRO
; APPLICANT: SEKI, NAOHICO
; APPLICANT: YOSHIKAWA, TSUTOMU
; APPLICANT: OTSUKI, MOTOKYU
; APPLICANT: NAGAHARI, KENJI
; APPLICANT: MASUHO, YASUHIKO
; TITLE OF INVENTION: NOVEL FULL-LENGTH CDNA
; FILE REFERENCE: 084335/0160
; CURRENT APPLICATION NUMBER: US/10/094,749
; CURRENT FILING DATE: 2002-03-12
; PRIOR APPLICATION NUMBER: 60/350,435
; PRIOR FILING DATE: 2002-01-24
; PRIOR APPLICATION NUMBER: JP 2001-328381
; PRIOR FILING DATE: 2001-09-14
; NUMBER OF SEQ ID NOS: 3381
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 2500
; LENGTH: 554
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-094-749-2500

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Query Match	29.7%	Score	906	DB	12	Length	554
Best Local Similarity	39.2%	Pred.	No. 5.6e-76				
Matches	208	Conservative	89	Mismatches	177	Indels	56
Gaps	12						
Qy	36	EAEALAPSGVWVPGARGGALARAAGRL	---	LALLLAVSAPRLQAEELGDGCGHLV	92		
Db	17	QVTRAAAAPAWAALPLSRSLPPCSN	SSSSPMFL	LLLLLVLLLLBEDAGQQDGGCHTV	76		
Qy	93	TYQDSGTWTSKNYPGTYPNHTVEK	ITVPGKRLILRLGDL	DIE-SQTCA	SYLLFTS-150		
Db	77	LGPESGTGLTSINYPQTVPNSTVC	EWELRIVQMGERVIRKFGD	FDIEDSDSCHFN	YLAIYNG136		
Qy	151	---SSDQGYDYG- SMTVPKELLNT	SEVTVPRESGSHSGRGFLTYA	SSDHPD	LIITCL206		
Db	137	IGVSRTEIGKYGCLQMHSIESK	NEITLLFPMGSIHVSGRFLAS	YVIDKQD	LIITCL196		
Qy	207	ERASHYKTEYSKFCPAGCRDVAGD	ISGNMVDGYRDTSL	LCKAAIHAGIIAD	BLGGQISV266		
Db	197	DTASNFLEPEFSKYCPAGCLLP	PAEISGTHPHGYRDSPLCMG	VGHAGVSVNTLGGQISV	256		
Qy	267	LQRKGSRYEGILANGVSRDGS	LSDKRFLFTSNGCSR	LSFE- ---PDQIRASS	SWQS322		
Db	257	VISKGIPYESSLANNVTSVGH	LSLSTLFTFKTSGCYGL	MESGVIA	DPOQTASSVLEW316		
Qy	323	VNESGQVHWSPGQARLQDQGS	WAGSDSNHKKPREWLEID	GEKKITGITRT	SGTQS382		
Db	317	TDHTGQENSWKPKARLKKP	GPWPAATDE- ---YQLQID	LNKEKKITGIT	TGSTMV372		
Qy	383	NFNFYUKSPFMNFKNNSKWK	TKYKGIWNNEEKVFGNS	NFRDPVQNNF	PIPIVARVRVU442		
Db	373	EHNYYSAYRILCSDGQKWTV	REPGVEQDKIFQGNKDY	HDVRN	NFLPIIAREFIRVN432		
Qy	443	PQWTHORIALKVELIGCQ	-----ITQ-----	GNDLSVWRKTSQ	STSVTKKEDT487		
Db	433	PTWOQOKIAMKVELLGCQ	IPKGRPPKLTQPPPP	NSND- ---LQNTAPP	KIAKGRAPK488		
Qy	488	ITRPI-----PSEET--	STGINITY-----	VAIPLV	LVLVIV157		
Db	489	FTOPLQPRSNPEAOTEOCTA	SPDIRNTTVPNTVKD	VALA	AVLPVLV338		

Search completed: December 29, 2003, 14:30:48
Job time : 646 secs

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Db	28	CGRLNSKDAGVITSGPYQDPYPSHQNCWVWYAPENQKIVLNFNPHFIEKHDCCKYDF	87
Qy	146	L4F-----TSSSQYQYPCGSMTPVKXELLNTSEVTVFESGSHISGRGFLTY-----AS	196
Db	88	IEIRDGSESADLLGHCHNI-APTIISSGSVYIKFTSDYARQAGFSLRYEIPKTS	146
Qy	197	SD-----HPDLITC-----LERASHVILKTSYXF-----	220
Db	147	EDCSKNFTSPNGTIBSPGPEKYPINLOCTTTIILAKPRMEILOFLTFDLEHDPLOVGEG	206
Qy	221	-CPAGCRDVAGDIS--GNMVDGYRDTSLCKAAIHAGIADLGGQISVLQRKISRYESG	277
Db	207	DKYDMLDITWDGIPHVGPLIGYCGTKPSKLRSTGILSLTFTHNAV-----	255
Qy	278	ILANGVLSRSGLSDKRFLFTSN-----GCSRSLSFEP---DQOIRASSWSQVNESGD	328
Db	256	-----AKDG-FSARYILVHQEPENFCQCAPLGMESGRIVNEQISASSTF-----S	300
Qy	329	QVHWSFGQARLOQGFPSWASGDSSNNHKPREWLIEDLGKKKITGRTTG--STQSNFNF	386
Db	301	DGRWTFQQSKRLHGGDDNGWTFPVDSN- ---KEYLOVDLRFLTMLTAITAQSAISRETQGY	356
Qy	387	YKYSFVWFNFNNNSKWTKYGLVINEEKVFCQNSNFRDPVQNNFPIPIVARYVVRVPQTW	446
Db	357	YKYSYKLEVSTNGEDWVYRHGKNH--KVFOANNDATEVLNKLHTPLLTRFIRIRPQTW	414
Qy	447	HQBIALKVELICQITQGNDSLWVRKTS-----QSTSVSTKKEDEITRP	491
Db	415	HLCIALRLLEFGCRVTDAPCSNMLGMSGLIADTOISASSTREYLWSP	462

RESULT 2

```

US-08-936-135-18
; Sequence 18, Application US/08936135
; Patent No. 6054293
; GENERAL INFORMATION:
; APPLICANT: Tessier-Lavigne, Marc
; APPLICANT: He, Zhigang
; APPLICANT: Chen, Hang
; TITLE OF INVENTION: Semaphorin Receptors
; NUMBER OF SEQUENCES: 26
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: SCIENCE & TECHNOLOGY LAW GROUP
; STREET: 75 DENISE DRIVE
; CITY: HILLSBOROUGH
; STATE: CALIFORNIA
; COUNTRY: USA
; ZIP: 94010
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/936,135
; FILING DATE:
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: OSMAN, RICHARD A
; REGISTRATION NUMBER: 36,627
; REFERENCE/DOCKET NUMBER: UC97-288-2
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (650) 343-4341
; TELEFAX: (650) 343-4342
; INFORMATION FOR SEQ ID NO: 18:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 909 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-936-135-18

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Query Match	11.4%; Score 349; DB 3; Length 909;
Best Local Similarity	25.3%; Pred. No. 1.4e-23;
Matches 109; Conservative	72; Mismatches 166; Indels 84; Gaps 16

QY	88	CGHLVTQDSGTMTSKNPGYTNHTVCEKTIITVPK-GKRLILRLG-DLDIESQTCASDY 145
DB	28	CGGRLSKSDAGYIITSPGYPODPYSHQNCCEWIVYAPENQKIVLNFNPHFEIEKHDKYDF 87
QY	146	LLF-----TSSDDQGYPCGSMVTPKELLNLTSEVTRFESGSHISGRGFLITY-----AS 196
DB	88	IEIRDGUSEADLLGKHGONI-APTIISSGSMLYIKFTSDYAROGAGFSRLYEIPKTCG 146
QY	197	SD-----HPDLITC-----LERASHYLKTEYSKF----- 220
DB	147	EDCSKNFTSPNGTIESPGFPEKYPHNLDOCTFIIAKPKWEIILQFLIFDLEHDPQLQVGBG 206
QY	221	-CPAGCRDVAGDIS--GNMVDGYRDTSLCKAAIHAGIIABELGQISVLQKRGISRVBG 277
DB	207	DKRYDWLIDWPIGHVUGPLIGKYCGTKTPSELRSSTGILSTLFTHTDMAVAK-----DG 259
QY	278	ILANGVLSRDCGSLDKRFLFTSNGCSRSLSPF---PDGQIRASSWSQSVNESGDQVHWS 333
DB	260	FSARYLHVQEPLENFQ-----CNVPLGHESGRINEQISASTY-----SDGRWT 305
QY	334	PGQARLQDQGSWASGSDSNHKNHPREWLEIDLGEKKKITGIRTTG--STQSNFNFYVKSF 391
DB	306	PQQRSLHGDDNGWTPNLDSN---KEYLQVDLRFULTMLTAITATQGAISRETQNGYVVKSY 361
QY	392	VMTFNKNNSKWTKYKGIYVNNKEKVPQGSNFRDPVQNNFPIPIVARYVRVVDQTHQRIA 451
DB	362	KLEVSTNGEDMMVYRHGKNH--KVPQANNDATEVVLNKLHAPLLTRFVRIRPQTWHSGIA 419
QY	452	LKVELIGCOIT 462
DB	420	LRLEFGCRVT 430

RESULT 3

US-08-936-135-20

; Sequence 20, Application US/08936135

; Patent No. 6054293

; GENERAL INFORMATION:

; APPLICANT: Tessier-Lavigne, Marc

; APPLICANT: He, Zhigang

; APPLICANT: Chen, Hang

; TITLE OF INVENTION: Semaphorin Receptors

; NUMBER OF SEQUENCES: 26

; CORRESPONDENCE ADDRESS:

; ADDRESSEE: SCIENCE & TECHNOLOGY LAW GROUP

; STREET: 75 DENISE DRIVE

; CITY: HILLSBOROUGH

; STATE: CALIFORNIA

; COUNTRY: USA

; ZIP: 94010

; COMPUTER READABLE FORM:

; MEDIUM TYPE: Floppy disk

; COMPUTER: IBM PC compatible

; OPERATING SYSTEM: PC-DOS/MS-DOS

; SOFTWARE: PatentIn Release #1.0, Version #1.30

; CURRENT APPLICATION DATA:

; APPLICATION NUMBER: US/08/936,135

; FILING DATE:

; CLASSIFICATION: 435

; ATTORNEY/AGENT INFORMATION:

; NAME: OSMAN, RICHARD A

; REGISTRATION NUMBER: 36,627

; REFERENCE/DOCKET NUMBER: UC97-288-2

; TELECOMMUNICATION INFORMATION:

; TELEPHONE: (650) 343-4341

; TELEFAX: (650) 343-4342

; INFORMATION FOR SEQ ID NO: 20:

; SEQUENCE CHARACTERISTICS:

; LENGTH: 926 amino acids

```
;
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-936-135-20

Query Match 11.4%; Score 349; DB 3; Length 926;
Best Local Similarity 25.3%; Pred. No. 1.4e-23;
Matches 109; Conservative 72; Mismatches 166; Indels 84; Gaps 16;

Qy 88 CGHLVTVQDSQGTMTSKNYPGTYPNHTVCEKTIITVPK-GKRLILRLG-DLDIESQTCASDY 145
Db 28 CGRLNSKAGYITSPGYPODYPSHQNCENVIVAPENQKIVLNFNPHFEIEKHDKYDF 87
Qy 146 LLF-----TSSSDQGYPCGSMVTPKELLNTSEVTVRFESGSHISGRGFLITY-----AS 196
Db 88 IEIRDGSESADLLGKHCGNI-APPTIISSGSMLYIKFTSDYARQAGFSLRYEIPKTS 146
Qy 197 SD-----HPDLITC-----LERASHYLKTEYSKP----- 220
Db 147 EDCSKNFTSPNGTIESPGPEKYPHNLDCFTTILAKPKMEIILQFLIFDLEHDPLOVGEG 206
Qy 221 -CPAGCRDVAGDIS--GNMVDGYRDTSLCKAAIHAGIHADELGGQISVLQKGISRYEG 277
Db 207 DCKYDWDLDWDGPHVGLIGKYCGTKTPSELRSSTGLSLTFHTDMAVAK-----DG 259
Qy 278 ILANGVLSRDGSLDKRFLFTSNGCSRSLSGPE-----PDGQIRASSWSQSVNESGDQVHWS 333
Db 260 FSARYYLHVOELENFO-----CNVPLGNESGRIANEQISASSTV-----SDGRWT 305
Qy 334 PGQARLQDQGPSWASGDSNNHKPREWLEIDLGEKKKITGIRTTG--STOSNFNFYVKS 391
Db 306 PQQSRLHGGDNGWTPNLDN-----KEYLQVDLRLTMTLTAITATQGAISRETQNGYVKS 361
Qy 392 VMKFNKNSKWKTKYGIWNEEKVFGNSFRDPVQNNFPIPVARYVRVVPQTHORIA 451
Db 362 KLEVSTNGEDMMVYRHGKNH--KVFOANNDATEVVLNKLHAPLTLTRFVRIRPQTHSGIA 419
Qy 452 LKVELIGCOIT 462
Db 420 LRLELFCRV 430

RESULT 4
US-08-936-135-22
; Sequence 22, Application US/08936135
; Patent No. 6054293
; GENERAL INFORMATION:
; APPLICANT: Tessier-Lavigne, Marc
; APPLICANT: He, Zhigang
; APPLICANT: Chen, Hang
; TITLE OF INVENTION: Semaphorin Receptors
; NUMBER OF SEQUENCES: 26
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: SCIENCE & TECHNOLOGY LAW GROUP
; STREET: 75 DENISE DRIVE
; CITY: HILLSBOROUGH
; STATE: CALIFORNIA
; COUNTRY: USA
; ZIP: 94010
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/936,135
; FILING DATE:
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: OSMAN, RICHARD A
; REGISTRATION NUMBER: 36, 627
; REFERENCE/DOCKET NUMBER: UC97-288-2

;
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (650) 343-4341
; TELEFAX: (650) 343-4342
; INFORMATION FOR SEQ ID NO: 22:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 901 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-936-135-22

Query Match 11.3%; Score 343.5; DB 3; Length 901;
Best Local Similarity 23.2%; Pred. No. 4.4e-23;
Matches 111; Conservative 72; Mismatches 177; Indels 119; Gaps 15;

Qy 88 CGHLVTVQDSQGTMTSKNYPGTYPNHTVCEKTIITVPK-GKRLILRLG-DLDIESQTCASDY 145
Db 28 CGRPNKAGYITSPGYPODYPSHQNCENVIVAPENQKIVLNFNPHFEIEKHDKYDF 87
Qy 146 LLF-----TSSSDQGYPCGSMVTPKELLNTSEVTVRFESGSHISGRGFLITY-----AS 196
Db 88 IEIRDGSESADLLGKHCGNI-APPTIISSGSMVLYIKFTSDYARQAGFSLRYEIPKTS 146
Qy 197 SD-----HPDLITC-----LERASHYLKTEYSKP----- 220
Db 147 EDCSKNFTSPNGTIESPGPEKYPHNLDCFTTILAKPKMEIILQFLIFDLEHDPLOVGEG 206
Qy 221 -CPAGCRDVAGDIS--GNMVDGYRDTSLCKAAIHAGIHADELGGQISVLQKGISRY-- 275
Db 207 DCKYDWDLDWDGPHVGLIGKYCGTKTPSELRSSTGLSLTFHTDMAVAKDGFSAARYL 266
Qy 276 -----EGILANGVLSRDGSLDKRFLFTSNGCSRSLSFEPDQGIARAS 317
Db 267 IHQEPENFOCNVPLGNESGRIANEQISASSTFSDGR----- 303
Qy 318 SSMQSVNESGDQVHWSFGQARLQDQGPSWASGDSNNHKPREWLEIDLGEKKKITGIRTT 377
Db 304 -----WTPQQSRLHGGDNGWTPNLDN-----KEYLQVDLRLTMTLTAITATQ 345
Qy 378 G--STOSNFNFYVKSFMKFNKNSKWKTKYGIWNEEKVFGNSFRDPVQNNFPIPV 435
Db 346 GAISRETQNGYVKSYLEVSTNGEDMMVYRHGKNH--KIFOANNDATEVVLNKLHAPL 403
Qy 436 ARYVRVVPQTHORIAKVELIGCOITQGNDSLVWRKTS---QSTSVSTKKEDETI 491
Db 404 TRFIRPQTHLGLIALRLLEFCRVTDAPCSNMLGSLIADTQISASSTREYLMSP 462

RESULT 5
US-08-936-135-24
; Sequence 24, Application US/08936135
; Patent No. 6054293
; GENERAL INFORMATION:
; APPLICANT: Tessier-Lavigne, Marc
; APPLICANT: He, Zhigang
; APPLICANT: Chen, Hang
; TITLE OF INVENTION: Semaphorin Receptors
; NUMBER OF SEQUENCES: 26
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: SCIENCE & TECHNOLOGY LAW GROUP
; STREET: 75 DENISE DRIVE
; CITY: HILLSBOROUGH
; STATE: CALIFORNIA
; COUNTRY: USA
; ZIP: 94010
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/936,135
```

```
;
; FILING DATE:
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: OSMAN, RICHARD A
; REGISTRATION NUMBER: 36,627
; REFERENCE/DOCKET NUMBER: UC97-288-2
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (650) 343-4341
; TELEFAX: (650) 343-4342
; INFORMATION FOR SEQ ID NO: 24:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 906 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; US-08-936-135-24

Query Match 11.3%; Score 343.5; DB 3; Length 906;
Best Local Similarity 23.2%; Pred. No. 4.5e-23;
Matches 111; Conservative 72; Mismatches 177; Indels 119; Gaps 15;

Qy 88 CGHLVTYQDSGWTSMKNYPGTYPNHTVCBKTITVPK-GKRLILRLG-DLDIESQTCASY 145
Db 28 CGGRPNKADAGYITSPGYPDYPSHQNCWEIVYAPEPNQKIVLNFNPFIEKHDKYDF 87
Qy 146 LLF-----TSSSDQYGPYCGSMVTPKELLNTSEVTVRFESGSHISGRGFLTY-----AS 196
Db 88 IEIRDGSESADLLGKHGNI-APPTIISSGSVLYIKFTSDYARQAGFSLRYEIKFTGS 146
Qy 197 SD-----HPDLITC-----LERASHYLKTEYSKF----- 220
Db 147 EDCSKNFTSPNGTIESPGFPEKYPHNLDCTFTILAKPRMEIILQFLTFDLEHDPLOVGE 206
Qy 221 -CPAGCRDVAGDIS--GNMVDGYRDTSLCKAAIHAGIIADELGGQISVLQRKISRY-- 275
Db 207 DCKYDMLDIWDGIPHVGLIKYCGTKPESKLSRSTGILSLTFHTDMAVAKDGFSAARYL 266
Qy 276 -----EGILANGVLSRDGSLSDKRFLLTSNGCSRSLSFEPDGOIRAS 317
Db 267 IHQEPPEFQCNVPLGMESGRIANEQISASTFSDGR----- 303
Qy 318 SSMQSVNESGDQVHWSFGQARLQDQGPSWASGDSNNHNPREWLEIDLGEKKKITGIRTT 377
Db 304 -----WTFQSRHLHGDGNGWTPNLDN-----KEYLQVDLRLTMTAIAATQ 345
Qy 378 G--STQSNFNFYKSFVWNFKNNSKWTKYKGIIVNNEEKVFGNSFRDPVQNNFPIPIV 435
Db 346 GAISRETQKGYVYKSKLEVSTNGEDWMVYRHGKNH--KIFQANNDATVYVNLKLMPL 403
Qy 436 ARYVRVVPOTWHORIALKVELIGCOITQGNDSLVRKTS---QSTSVSTKKEDETTIRP 491
Db 404 TRFIRIRPQTHLGIARLELFGCRVTDAPCSNMLGMLSLGIADTQISASTREYLWSP 462

RESULT 6
US-08-936-135-8
; Sequence 8, Application US/08936135
; Patent No. 6054293
; GENERAL INFORMATION:
; APPLICANT: Tessier-Lavigne, Marc
; APPLICANT: He, Zhigang
; APPLICANT: Chen, Hang
; TITLE OF INVENTION: Semaphorin Receptors
; NUMBER OF SEQUENCES: 26
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: SCIENCE & TECHNOLOGY LAW GROUP
; STREET: 75 DENISE DRIVE
; CITY: HILLSBOROUGH
; STATE: CALIFORNIA
; COUNTRY: USA
; ZIP: 94010
; COMPUTER READABLE FORM:

;
; FILING DATE:
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: OSMAN, RICHARD A
; REGISTRATION NUMBER: 36,627
; REFERENCE/DOCKET NUMBER: UC97-288-2
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (650) 343-4341
; TELEFAX: (650) 343-4342
; INFORMATION FOR SEQ ID NO: 24:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 906 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; US-08-936-135-24

Query Match 11.3%; Score 343.5; DB 3; Length 906;
Best Local Similarity 23.2%; Pred. No. 4.5e-23;
Matches 111; Conservative 72; Mismatches 177; Indels 119; Gaps 15;

Qy 88 CGHLVTYQDSGWTSMKNYPGTYPNHTVCBKTITVPK-GKRLILRLG-DLDIESQTCASY 145
Db 28 CGGRPNKADAGYITSPGYPDYPSHQNCWEIVYAPEPNQKIVLNFNPFIEKHDKYDF 87
Qy 146 LLF-----TSSSDQYGPYCGSMVTPKELLNTSEVTVRFESGSHISGRGFLTY-----AS 196
Db 88 IEIRDGSESADLLGKHGNI-APPTIISSGSVLYIKFTSDYARQAGFSLRYEIKFTGS 146
Qy 197 SD-----HPDLITC-----LERASHYLKTEYSKF----- 220
Db 147 EDCSKNFTSPNGTIESPGFPEKYPHNLDCTFTILAKPRMEIILQFLTFDLEHDPLOVGE 206
Qy 221 -CPAGCRDVAGDIS--GNMVDGYRDTSLCKAAIHAGIIADELGGQISVLQRKISRY-- 275
Db 207 DCKYDMLDIWDGIPHVGLIKYCGTKPESKLSRSTGILSLTFHTDMAVAKDGFSAARYL 266
Qy 276 -----EGILANGVLSRDGSLSDKRFLLTSNGCSRSLSFEPDGOIRAS 317
Db 267 IHQEPPEFQCNVPLGMESGRIANEQISASTFSDGR----- 303
Qy 318 SSMQSVNESGDQVHWSFGQARLQDQGPSWASGDSNNHNPREWLEIDLGEKKKITGIRTT 377
Db 304 -----WTFQSRHLHGDGNGWTPNLDN-----KEYLQVDLRLTMTAIAATQ 345
Qy 378 G--STQSNFNFYKSFVWNFKNNSKWTKYKGIIVNNEEKVFGNSFRDPVQNNFPIPIV 435
Db 346 GAISRETQKGYVYKSKLEVSTNGEDWMVYRHGKNH--KIFQANNDATVYVNLKLMPL 403
Qy 436 ARYVRVVPOTWHORIALKVELIGCOITQGNDSLVRKTS---QSTSVSTKKEDETTIRP 491
Db 404 TRFIRIRPQTHLGIARLELFGCRVTDAPCSNMLGMLSLGIADTQISASTREYLWSP 462

RESULT 7
US-08-936-135-10
; Sequence 10, Application US/08936135
; Patent No. 6054293
; GENERAL INFORMATION:
; APPLICANT: Tessier-Lavigne, Marc
; APPLICANT: He, Zhigang
; APPLICANT: Chen, Hang
; TITLE OF INVENTION: Semaphorin Receptors
; NUMBER OF SEQUENCES: 26
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: SCIENCE & TECHNOLOGY LAW GROUP
```

STREET: 75 DENISE DRIVE
CITY: HILLSBOROUGH
STATE: CALIFORNIA
COUNTRY: USA
ZIP: 94010
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent In Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/936,135
FILING DATE:
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: OSMAN, RICHARD A
REGISTRATION NUMBER: 36,627
REFERENCE/DOCKET NUMBER: UC97-288-2
TELEPHONE: (650) 343-4341
TELEFAX: (650) 343-4342
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 909 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-936-135-10

Query Match 11.3%, Score 343.5; DB 3; Length 909;
Best Local Similarity 23.2%; Pred. No. 4.5e-23;
Matches 111; Conservative 72; Mismatches 177; Indels 119; Gaps 15;
QY 88 CGHLVTVQDSGTMKPNYPCGYTHVCEKTIIVPK-GKRLILRLG-DLIESQTCASDY 145
Db 28 CGRPNKSDAGYITSPGYDPYHQNCWEIVVAPENQKIVLNFNPFIEKHDKYDF 87
QY 146 LLF-----TSSSDQVPGYCGSMVTPKELLNTSEVTVRFSGSHISGRGFLTY-----AS 196
Db 88 IEIRDGSESADLLGKHGNI-APPTIISSGSVLYIKFTSDYARQAGFSLRVEIFKTS 146
QY 197 SD-----HPDLITC-----LERASHYLKTEYSKF----- 220
Db 147 EDCSKNFTSPNGTIESPGPEKYPHNLDCTFTILAKPRMEIILQFLTFDLEHDPQVGE 206
QY 221 -CPAGCEDVAGDIS--GNMVDGYRDTSLCKAAIHAGIIADELGGQISVLQKGISY-- 275
Db 207 DCKYDWDLDWDGIPHVGLIGKYCGTKTPSKLSRSTGILSLFTHTDMAVAKDGFSA 266
QY 276 -----EGILANGVLSRDGSLSDKRFLLTSPNGCSRSLSFEPDQGI 317
Db 267 IQEPPENFCNVPLMGESGRIANEQISASTFSDGR----- 303
QY 318 SSWQSVNESGDQVHWSFGQARLQDQGPSWASGDSNNHKKPREWLEIDLGEKKKITGIRTT 377
Db 304 -----WTPQQRSLHGGDNGWTPLNDSN-----KEYLQVDLRLFTMLTATQ 345
QY 378 G--STQSNFNFYKSFVMNFKNNKWKTKYKGIIVNNEEKVFGNSNFRDPVQNNFPIV 435
Db 346 GAISRETQKGYVYKSKLEVSTNGEDWMVYRHGKNH--KIFQANNDATEVVLNKLHMP 403
QY 436 ARYVRVVPQTHQRIALVELIGCQITQGNDSLVRKTS---QSTSVSTKKEDETI 491
Db 404 TRFIRPQTHLGLIALRLLELFGCRVTDAPCSNMLGSLGIADTQISASSTREYLWSP 462

RESULT 8

US-08-936-135-12
Sequence 12, Application US/08936135
Patent No. 6054293
GENERAL INFORMATION:
APPLICANT: Tessier-Lavigne, Marc

APPLICANT: He, Zhigang
APPLICANT: Chen, Hang
TITLE OF INVENTION: Semaphorin Receptors
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: SCIENCE & TECHNOLOGY LAW GROUP
STREET: 75 DENISE DRIVE
CITY: HILLSBOROUGH
STATE: CALIFORNIA
COUNTRY: USA
ZIP: 94010
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent In Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/936,135
FILING DATE:
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: OSMAN, RICHARD A
REGISTRATION NUMBER: 36,627
REFERENCE/DOCKET NUMBER: UC97-288-2
TELEPHONE: (650) 343-4341
TELEFAX: (650) 343-4342
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
LENGTH: 914 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-936-135-12

Query Match 11.3%, Score 343.5; DB 3; Length 914;
Best Local Similarity 23.2%; Pred. No. 4.5e-23;
Matches 111; Conservative 72; Mismatches 177; Indels 119; Gaps 15;
QY 88 CGHLVTVQDSGTMKPNYPCGYTHVCEKTIIVPK-GKRLILRLG-DLIESQTCASDY 145
Db 28 CGRPNKSDAGYITSPGYDPYHQNCWEIVVAPENQKIVLNFNPFIEKHDKYDF 87
QY 146 LLF-----TSSSDQVPGYCGSMVTPKELLNTSEVTVRFSGSHISGRGFLTY-----AS 196
Db 88 IEIRDGSESADLLGKHGNI-APPTIISSGSVLYIKFTSDYARQAGFSLRVEIFKTS 146
QY 197 SD-----HPDLITC-----LERASHYLKTEYSKF----- 220
Db 147 EDCSKNFTSPNGTIESPGPEKYPHNLDCTFTILAKPRMEIILQFLTFDLEHDPQVGE 206
QY 221 -CPAGCEDVAGDIS--GNMVDGYRDTSLCKAAIHAGIIADELGGQISVLQKGISY-- 275
Db 207 DCKYDWDLDWDGIPHVGLIGKYCGTKTPSKLSRSTGILSLFTHTDMAVAKDGFSA 266
QY 276 -----EGILANGVLSRDGSLSDKRFLLTSPNGCSRSLSFEPDQGI 317
Db 267 IQEPPENFCNVPLMGESGRIANEQISASTFSDGR----- 303
QY 318 SSWQSVNESGDQVHWSFGQARLQDQGPSWASGDSNNHKKPREWLEIDLGEKKKITGIRTT 377
Db 304 -----WTPQQRSLHGGDNGWTPLNDSN-----KEYLQVDLRLFTMLTATQ 345
QY 378 G--STQSNFNFYKSFVMNFKNNKWKTKYKGIIVNNEEKVFGNSNFRDPVQNNFPIV 435
Db 346 GAISRETQKGYVYKSKLEVSTNGEDWMVYRHGKNH--KIFQANNDATEVVLNKLHMP 403
QY 436 ARYVRVVPQTHQRIALVELIGCQITQGNDSLVRKTS---QSTSVSTKKEDETI 491
Db 404 TRFIRPQTHLGLIALRLLELFGCRVTDAPCSNMLGSLGIADTQISASSTREYLWSP 462

Db 304 -----WTQQSLHGDGNGWTFNLSN-----KEYLQVDRFLTMLTAIATQ 345
Qy 378 G--STQSNFNFYKSFVNFNNKWKTYKGIIVNNEEKVFGNSNFRDPVQNNFIPPIV 435
Db 346 GALSIRETKQYVYKYLEVSTNGEDMMVYRHGKNH--KIFQANNDATEVVLNKLHMLL 403
Qy 436 ARYRVVVPQTHORIALKVELIGCOITQGNDSLVRKTS---QSTSVSTKKEDETTIRP 491
Db 404 TRFIRIPQTHLGLALRLFLGCRVTDAPCSNMLGSLGLIADTQISASSTREYLWSP 462

RESULT 11
US-09-116-473-4
; Sequence 4, Application US/09116473
; Patent No. 6428965
; GENERAL INFORMATION:
; APPLICANT: Kolodkin, Alex
; APPLICANT: Ginty, David
; TITLE OF INVENTION: SEMAPHORIN RECEPTOR
; NUMBER OF SEQUENCES: 8
; CORRESPONDENCE ADDRESSES:
; ADDRESSEE: Banner & Witcoff
; STREET: 1001 G Street, NW
; CITY: Washington
; STATE: DC
; COUNTRY: USA
; ZIP: 20001
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: DOS
; SOFTWARE: FastSeq for Windows Version 2.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/116,473
; FILING DATE: 17-JUL-1998
; CLASSIFICATION:
; PRIORITY APPLICATION DATA:
; APPLICATION NUMBER: 60/052,762
; FILING DATE: 17-JUL-1998
; ATTORNEY/AGENT INFORMATION:
; NAME: Kagan, Sarah A
; REGISTRATION NUMBER: 32141
; REFERENCE/DOCKET NUMBER: 01107.74973
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 202-508-9100
; TELEFAX: 202-508-9299
; TELEX:
; INFORMATION FOR SEQ ID NO: 4:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 922 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; US-09-116-473-4

Query Match 10.8%; Score 331; DB 4; Length 922;
Best Local Similarity 24.1%; Pred. No. 6.9e-22;
Matches 112; Conservative 81; Mismatches 166; Indels 106; Gaps 19;

Qy 64 RGL-----LALLVAVSAPRLQAEELGDCGHLVTVYQDSGTMTSKNYPGTYPNHTVCEK 117
Db 3 RGLPLLCATLALALAGAFR-----SDKCGGTIKIENPGYLTPGYPHSHYHPSEKCEW 56
Qy 118 TITVPKG-KRLIIRLG-DLDIESQTCASDYLLFTSSDQ-----YGPYCGSMTVPKELLN 171
Db 57 LIQAPEYQRIIMFNPHFPLEDRDCKYDVEIDGENEGRLWGKFCGKI-APSPVSS 115
Qy 172 TSEVTRFEGSGSHISGRGFLTYA-----SSDHPDLITC-- 205
Db 116 GPFLFKFVSDYETHGAGFSIRIEIFKRGPECSONYTAFTGVKISPGFPEKYPNSLECTY 175
Qy 206 -----LERASHYLKTEYSKFCPAG--CRDVAGDISGNMVDGYRDTSLCKAAIH 252

Db 176 IIFAPKMSSEILBFESFDLEQDSNP--PGVFCRYDRLEI-----WDGFPE----- 219
Qy 253 AGIADLGLGQISVLQKRGISRYEGILA-----NGVLSRDGSLDKRFLFTSNG-----CS 303
Db 220 ---VGPHIGREYCCQKTPGRIRSSSGILSMVFYTDALAKEGFSANYSVLOSSISEDPKCM 276
Qy 304 RSLSPFP---DCQIRASSSWQSVNBSGDQVHWSQOARLODQGPSWASGDSSNNHKKPRE 359
Db 277 EALGMESGEIHSQITASSQYGT-----NMSVERSLNYPENGWTPGEDSY-----RE 324
Qy 360 WLEIDLGEKKKITGRTTG--STQSNFNFYKSFVNFNNKWKTYKGIIVNNEEKVQ 417
Db 325 WIQVDLGLLRFVTAAGTQGAISKETKKYKYYKTYRVDISSNGEDWTLK--EGNKALIFQ 382
Qy 418 GNSNFRDPVQNNFIPPIVARYVRVWPQTWHORIALKVELIGCOIT 462
Db 383 GNTNPTDVVFGVPFKPLITRFVRIKASWETGISMRFVYGCKIT 427

RESULT 12
US-08-936-135-6
; Sequence 6, Application US/08936135
; Patent No. 6054293
; GENERAL INFORMATION:
; APPLICANT: Tessier-Lavigne, Marc
; APPLICANT: He, Zhigang
; APPLICANT: Chen, Hang
; TITLE OF INVENTION: Semaphorin Receptors
; NUMBER OF SEQUENCES: 26
; CORRESPONDENCE ADDRESSES:
; ADDRESSEE: SCIENCE & TECHNOLOGY LAW GROUP
; STREET: 75 DENISE DRIVE
; CITY: HILLSBOROUGH
; STATE: CALIFORNIA
; COUNTRY: USA
; ZIP: 94010
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/936,135
; FILING DATE:
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: OSMAN, RICHARD A
; REGISTRATION NUMBER: 36,627
; REFERENCE/DOCKET NUMBER: UC97-288-2
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (650) 343-4341
; TELEFAX: (650) 343-4342
; INFORMATION FOR SEQ ID NO: 6:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 923 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; US-08-936-135-6

Query Match 10.8%; Score 329; DB 3; Length 923;
Best Local Similarity 24.0%; Pred. No. 1.1e-21;
Matches 113; Conservative 79; Mismatches 161; Indels 118; Gaps 21;

Qy 64 RGL-----LALLVAVSAPRLQAEELGDCGHLVTVYQDSGTMTSKNYPGTYPNHTVCEK 117
Db 3 RGLPLLCATLALALAGAFR-----SDKCGGTIKIENPGYLTPGYPHSHYHPSEKCEW 56
Qy 118 TITVPKG-KRLIIRLG-DLDIESQTCASDYLLFTSSDQ-----YGPYCGSMTVPKELLN 171
Db 57 LIQAPEYQRIIMFNPHFPLEDRDCKYDVEIDGENEGRLWGKFCGKI-APSPVSS 115

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Qy 172 TSEVTVFESGSHISGRGFLITVA-----SSDHPDLITC-- 205
Db 116 GPFLFIKFDVSDYETHGAGFSIRYEIFKRGPECSQNYTAPTGVIKSPGPFPEKYPNCLECTY 175
Qy 206 -----LERASHYLKTEYSKFCPAG--CRDVAGDISGNNVDGYRDTSLCKAAIH 252
Db 176 IIFAPKMSIILEPESPDLEQDSNP--PGGMFCRYDRLEI-----WDGFPE-----VGPH 223
Qy 253 AG-IIADELGGQI-----SVLORKGISRYEGILANGVLSDGSLSDKXFLF 297
Db 224 IGRYCGQKTPGRIRSSGVLMSWPFYDTSATAKEGFSANYSVLQSSI-SEDFK----- 274
Qy 298 TSNCSRSLSSEP-----DGOIRASSWSQSVNESGDQVHWSGQARLQDQGPSWASGDSSN 353
Db 275 -----CMEALGNESGEIHSDDQITASSQYGT-----NWSVERSLNYPENGWTPGEDSY 322
Qy 354 NHKPREMLEIDLGEKKKITGIRTTG--STQSNFNFYKSFVMNPKNNNSKWKTKYGVINN 411
Db 323 ----KEWIQVDLGLLRFTVAGTQGAISKETKKKYVKTVRVDISSNGEDWISLK--EGN 376
Qy 412 EEKVFQGSNFRDPVQNNFPIPIVARYVUVVPTWHQRIALKVELIGCOIT 462
Db 377 KALIFQGNTPDVTDLVGLVFSKPLITRFVRIKPVSWETGISMRFVYCGKIT 427

RESULT 13
US-07-607-538C-4
; Sequence 4, Application US/07607538C
; Patent No. 5455031
; GENERAL INFORMATION:
; APPLICANT: Ceriani Dr., Roberto L.
; APPLICANT: Peterson Dr., Jerry A.
; APPLICANT: Larocca, David J.
; TITLE OF INVENTION: POLYPEPTIDE WITH 46
; TITLE OF INVENTION: DIFFERENTIATION ANTIGEN BINDING SPECIFICITY AND CLOTTING
; TITLE OF INVENTION: FACTORS V AND VIII LIGHT-CHAIN HOMOLOGIES,
; TITLE OF INVENTION: FUSION PROTEIN, POLYNUCLEOTIDE AND POLYRIBO-
; TITLE OF INVENTION: NUCLEOTIDE ENCODING THE POLYPEPTIDE, ANTI-
; TITLE OF INVENTION: POLYPEPTIDE ANTIBODIES, KITS AND METHODS OF
; TITLE OF INVENTION: USE THEREOF
; NUMBER OF SEQUENCES: 5
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: V. Amzel & Assoc.
; STREET: 2055 No. 5455031th Broadway
; CITY: Walnut Creek
; STATE: California
; COUNTRY: USA
; ZIP: 94596
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS 5.0
; SOFTWARE: Patent In Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/07/607,538C
; FILING DATE: 01-NOV-1990
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Viviana Amzel
; REGISTRATION NUMBER: 30,930
; REFERENCE/DOCKET NUMBER: CRFCC-004
; TELEPHONE: (510) 943-1931
; TELEFAX: (510) 943-1189
; TELEX: N.A.
; INFORMATION FOR SEQ ID NO: 4:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 218 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
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; FRAGMENT TYPE:
US-07-607-538C-4
Query Match 10.5%; Score 321; DB 1; Length 218;
Best Local Similarity 42.6%; Pred. No. 5.5e-22;
Matches 72; Conservative 30; Mismatches 51; Indels 16; Gaps 5;
Qy 300 NGCSRSLSFE-----PDQQRASS---SQSVNESGDQVHWSGQARLQDQGPSWASGDSS 352
Db 58 NGCSTPLGMENKIKENKQITASSFKLSW-----GD--YWEPPFARLNAQGRVNAWQAKA 110
Qy 353 NNHKPREMLEIDLGEKKKITGIRTTG--STQSNFNFYKSFVMNPKNNNSKWKTKYGVINNE 412
Db 111 NNNK--QWLEIDLKIKKTAITITQCKSLSSSEMYKSYTIHYSEQGVENKPYRLKSSMV 168
Qy 413 EEKVFQGSNFRDPVQNNFPIPIVARYVUVVPTWHQRIALKVELIGCOI 461
Db 169 DKIFEGNTNTKGHVKNQFNFNPPIISRFIRVTPKTNOSIALRLLELFGCDI 217

RESULT 14
US-08-162-402B-4
; Sequence 4, Application US/08162402B
; Patent No. 5972337
; GENERAL INFORMATION:
; APPLICANT: CERIANI, ROBERTO L.
; APPLICANT: PETERSON, JERRY A.
; APPLICANT: LARocca, DAVID J.
; TITLE OF INVENTION: 46 KDALTON HUMAN MILK FAT
; TITLE OF INVENTION: GLOBULE (HMFG) ANTIGEN, FRAGMENTS & FUSION PROTEIN
; NUMBER OF SEQUENCES: 29
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Pretty, Schroeder & Poplawski
; STREET: 444 South Flower St., 19th Floor
; CITY: Los Angeles
; STATE: CA
; COUNTRY: USA
; ZIP: 90071
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: DOS
; SOFTWARE: FastSeq for Windows Version 2.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/162,402B
; FILING DATE: 03-DEC-1993
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER:
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: Amzel, Viviana
; REGISTRATION NUMBER: 30,930
; REFERENCE/DOCKET NUMBER: P66 38215
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 213-622-7700
; TELEFAX: 213-489-4210
; TELEX:
; INFORMATION FOR SEQ ID NO: 4:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 218 amino acids
; TYPE: amino acid
; STRANDEDNESS: unknown
; TOPOLOGY: unknown
; MOLECULE TYPE: peptide
US-08-162-402B-4
Query Match 10.5%; Score 321; DB 2; Length 218;
Best Local Similarity 42.6%; Pred. No. 5.5e-22;
Matches 72; Conservative 30; Mismatches 51; Indels 16; Gaps 5;
Qy 300 NGCSRSLSFE-----PDQQRASS---SQSVNESGDQVHWSGQARLQDQGPSWASGDSS 352
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Db 58 NGCSTPLGMENKIQITASSFKKSW-----GD--YWEFPRARLNAGQVRNWAQAKA 110
QY 353 NNHKPREWLIDGEGKKITGIRTTGTSQSNFNFYKSFVMNFKNNSKWKTKYKGIYNNE 412
Db 111 NNNK--QWLEIDLKIKKTAITITQCKSLSSSEMYKYSIYHQSGVEMKPYRLKSSMV 168
QY 413 EKVFQGNPNRDPVQNNFPIPIVARYVRVPTWHQRIALKVELIGCQI 461
Db 169 DKIFEGNTTKGHVKNFNFNPPFIISRFIRIPIKTNQSIARLRLFLFGCDI 217

RESULT 15

US-08-746-111-5
; Sequence S, Application US/08746111
; Patent No. 6066778
; GENERAL INFORMATION:
; APPLICANT: Ginsburg, David
; APPLICANT: Cui, Jisong
; TITLE OF INVENTION: Compositions And Methods For Screening
; TITLE OF INVENTION: Compounds For Anticoagulant Activity
; NUMBER OF SEQUENCES: 54
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Medlen & Carroll, LLP
; STREET: 220 Montgomery Street, Suite 2200
; CITY: San Francisco
; STATE: California
; COUNTRY: United States of America
; ZIP: 94104
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent In Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/746,111
; FILING DATE: 06-NOV-1996
; CLASSIFICATION:
; ATTORNEY/AGENT INFORMATION:
; NAME: Ingolia, Diane E.
; REGISTRATION NUMBER: 40,027
; REFERENCE/DOCKET NUMBER: UM-02536
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (415) 705-8410
; TELEFAX: (415) 397-8338
; INFORMATION FOR SEQ ID NO: 5:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 2183 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-746-111-5

Query Match 10.4%; Score 318; DB 3; Length 2183;
Best Local Similarity 40.8%; Pred. No. 4.8e-20;
Matches 69; Conservative 31; Mismatches 53; Indels 16; Gaps 5;
QY 300 NGCSRSISSE-----PDGQIRASS---SQSVNESGDQVHMSGQARLQDQSPWASGDS 352
Db 2023 NGCSTPLGLEGRIDQKITASSFKKSW-----GD--YWEFPRARLNAGQVRNWAQAKA 2075
QY 353 NNHKPREWLIDGEGKKITGIRTTGTSQSNFNFYKSFVMNFKNNSKWKTKYKGIYNNE 412
Db 2076 NNNK--QWLQVLLKIKKTAITITQCKSLSSSEMYKYSIYHQSGVEMKPYRLKSSMV 2133
QY 413 EKVFQGNPNRDPVQNNFPIPIVARYVRVPTWHQRIALKVELIGCQI 461
Db 2134 DKIFEGNNTKGHMKNFNFNPPFIISRFIRIPIKTNQSIARLRLFLFGCDI 2182

Search completed: December 29, 2003, 14:34:30
Job time: 22 secs

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